

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
27 December 2001 (27.12.2001)

PCT

(10) International Publication Number
WO 01/97850 A2

- (51) International Patent Classification⁷: **A61K 45/06**
- (21) International Application Number: **PCT/EP01/06976**
- (22) International Filing Date: 20 June 2001 (20.06.2001)
- (25) Filing Language: English
- (26) Publication Language: English
- (30) Priority Data:
00250194.8 23 June 2000 (23.06.2000) EP
00250214.4 28 June 2000 (28.06.2000) EP
- (71) Applicant: **SCHERING AKTIENGESELLSCHAFT**
[DE/DE]; Müllerstrasse 178, 13353 Berlin (DE).
- (71) Applicants and
(72) Inventors: **SIEMEISTER, Gerhard** [DE/DE]; Reimerswalder Steig 26, 13503 Berlin (DE). **HABEREY, Martin** [DE/DE]; Steinstr. 1, 12169 Berlin (DE). **THIERAUCH, Karl-Heinz** [DE/DE]; Hochwildpfad 45, 14169 Berlin (DE).
- (81) Designated States (*national*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CR, CU, CZ, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW.
- (84) Designated States (*regional*): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

Published:

— without international search report and to be republished upon receipt of that report

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

WO 01/97850 A2

(54) Title: COMBINATIONS AND COMPOSITIONS WHICH INTERFERE WITH VEGF/VEGF AND ANGIOPOIETIN/TIE RECEPTOR FUNCTION AND THEIR USE (II)

(57) Abstract: The present invention describes the combination of substances interfering with the biological activity of Vascular Endothelial Growth Factor (VEGF)/VEGF receptor systems (compound I) and substances interfering with the biological function of Angiopoietin/Tie receptor systems (compound II) for inhibition of vascularization and for cancer treatment.

**Combinations and compositions which interfere with VEGF/ VEGF and
angiopoietin/ Tie receptor function and their use (II)**

- 5 The present invention provides the combination of substances interfering with the biological activity of Vascular Endothelial Growth Factor (VEGF)/VEGF receptor systems (compound I) and substances interfering with the biological function of Angiopoietin/Tie receptor systems (compound II) for inhibition of vascularization and for cancer treatment.
- 10 Protein ligands and receptor tyrosine kinases that specifically regulate endothelial cell function are substantially involved in physiological as well as in disease-related angiogenesis. These ligand/receptor systems include the Vascular Endothelial Growth Factor (VEGF) and the Angiopoietin (Ang) families, and their receptors, the VEGF receptor family and the tyrosine kinase with immunoglobulin-like and epidermal growth factor homology domains (Tie) family. The members of the two families of receptor tyrosine kinases are expressed primarily on endothelial cells. The VEGF receptor family includes Flt1 (VEGF-R1), Flk1/KDR (VEGF-R2), and Flt4 (VEGF-R3). These receptors are recognized by members of the VEGF-related growth factors in that the ligands of Flt1 are VEGF and placenta growth factor (PIGF), whereas Flk1/KDR binds VEGF, VEGF-C and VEGF-D, and the ligands of Flt4 are VEGF-C and VEGF-D (Nicosia, Am. J. Pathol. 153, 11-16, 1998). The second family of endothelial cell specific receptor tyrosine kinases is represented by Tie1 and Tie2 (also known as Tek). Whereas Tie1 remains an orphan receptor, three secreted glycoprotein ligands of Tie2, Ang1, Ang2, and Ang3/Ang4 have been discovered (Davis et al., Cell 87, 1161-1169, 1996; Maisonpierre et al., Science 277, 55-60, 1997; Valenzuela et al, Proc. Natl. Acad. Sci. USA 96, 1904-1909, 1999; patents: US 5,521,073; US 5,650,490; US 5,814,464).
- 15
- 20
- 25
- 30
- The pivotal role of VEGF and of its receptors during vascular development was exemplified in studies on targeted gene inactivation. Even the heterozygous disruption of the VEGF gene resulted in fatal deficiencies in vascularization (Carmeliet et al., Nature 380, 435-439, 1996; Ferrara et al., Nature 380, 439-442,

1996). Mice carrying homozygous disruptions in either Flt1 or Flk1/KDR gene die in mid-gestation of acute vascular defects. However, the phenotypes are distinct in that Flk1/KDR knock-out mice lack both endothelial cells and a developing hematopoietic system (Shalaby et al. *Nature* 376, 62-66, 1995), whereas Flt1 deficient mice have normal hematopoietic progenitors and endothelial cells, which fail to assemble into functional vessels (Fong et al., 376, 66-70, 1995). Disruption of the Flt4 gene, whose extensive embryonic expression becomes restricted to lymphatic vessels in adults, revealed an essential role of Flt4 for the remodeling and maturation of the primary vascular networks into larger blood vessels during early development of the cardiovascular system (Dumont et al., *Science* 282, 946-949, 1998). Consistent with the lymphatic expression of Flt4 in adults overexpression of VEGF-C in the skin of transgenic mice resulted in lymphatic, but not vascular, endothelial proliferation and vessel enlargement (Jeltsch et al., *Science* 276, 1423-1425, 1997). Moreover, VEGF-C was reported to induce neovascularization in mouse cornea and chicken embryo chorioallantoic membrane models of angiogenesis (Cao et al., *Proc. Natl. Acad. Sci. USA* 95, 14389-14394, 1998).

The second class of endothelial cell specific receptor tyrosine kinases has also been found to be critically involved in the formation and integrity of vasculature. Mice deficient in Tie1 die of edema and hemorrhage resulting from poor structural integrity of endothelial cells of the microvasculature (Sato et al., *Nature* 376, 70-74, 1995; Rodewald & Sato, *Oncogene* 12, 397-404, 1996). The Tie2 knock-out phenotype is characterized by immature vessels lacking branching networks and lacking periendothelial support cells (Sato et al., *Nature* 376, 70-74, 1995; Dumont et al., *Genes Dev.* 8, 1897-1909, 1994). Targeted inactivation of the Tie2 ligand Ang1, as well as overexpression of Ang2, an inhibitory ligand, resulted in phenotypes similar to the Tie2 knock out (Maisonpierre et al., *Science* 277, 55-60, 1997; Suri et al., *cell* 87, 1171-1180). Conversely, increased vascularization was observed upon transgenic overexpression of Ang1 (Suri et al., *Science* 282, 468-471, 1998; Thurstonen et al., *Science* 286, 2511-2514, 1999).

The results from angiogenic growth factor expression studies in corpus luteum development (Maisonpierre et al., *Science* 277, 55-60, 1997; Goede et al. Lab.

Invest. 78, 1385-1394, 1998), studies on blood vessel maturation in the retina (Alon et al., Nature Med. 1, 1024-1028, 1995; Benjamin et al, Development 125, 1591-1598, 1998), and gene targeting and transgenic experiments on Tie2, Ang1, and Ang2, suggest a fundamental role of the Angiopoietin/Tie receptor system in

- 5 mediating interactions between endothelial cells and surrounding pericytes or smooth muscle cells. Ang1, which is expressed by the periendothelial cells and seems to be expressed constitutively in the adult, is thought to stabilize existing mature vessels. Ang2, the natural antagonist of Ang1 which is expressed by endothelial cells at sites of vessel sprouting, seems to mediate loosening of
10 endothelial-periendothelial cell contacts to allow vascular remodeling and sprouting in cooperation with angiogenesis initiators such as VEGF, or vessel regression in the absence of VEGF (Hanahan, Science 277, 48-50, 1997).

- In pathological settings associated with aberrant neovascularization elevated expression of angiogenic growth factors and of their receptors has been observed.
15 Most solid tumors express high levels of VEGF and the VEGF receptors appear predominantly in endothelial cells of vessels surrounding or penetrating the malignant tissue (Plate et al., Cancer Res. 53, 5822-5827, 1993). Interference with the VEGF/VEGF receptor system by means of VEGF-neutralizing antibodies
20 (Kim et al., Nature 362, 841-844, 1993), retroviral expression of dominant negative VEGF receptor variants (Millauer et al., Nature 367, 576-579, 1994), recombinant VEGF-neutralizing receptor variants (Goldman et al., Proc. Natl. Acad. Sci. USA 95, 8795-8800, 1998), or small molecule inhibitors of VEGF receptor tyrosine kinase (Fong et al., Cancer Res. 59, 99-106, 1999; Wedge et al., Cancer Res. 60,
25 970-975, 2000; Wood et al. Cancer Res. 60, 2178-2189, 2000), or targeting cytotoxic agents via the VEGF/VEGF receptor system (Arora et al., Cancer Res. 59, 183-188, 1999; EP 0696456A2) resulted in reduced tumor growth and tumor vascularization. However, although many tumors were inhibited by interference with the VEGF/VEGF receptor system, others were unaffected (Millauer et al.,
30 Cancer Res. 56, 1615-1620, 1996). Human tumors as well as experimental tumor xenografts contain a large number of immature blood vessels that have not yet recruited periendothelial cells. The fraction of immature vessels is in the range of 40% in slow growing prostate cancer and 90% in fast growing glioblastoma. A selective obliteration of immature tumor vessels was observed upon withdrawal of

VEGF by means of downregulation of VEGF transgene expression in a C6 glioblastoma xenograft model. This result is in accordance with a function of VEGF as endothelial cell survival factor. Similarly, in human prostate cancer shutting off VEGF expression as a consequence of androgen-ablation therapy led to selective apoptotic death of endothelial cells in vessels lacking periendothelial cell coverage. In contrast, the fraction of vessels which resisted VEGF withdrawal showed periendothelial cell coverage (Benjamin et al., J. Clin. Invest. 103, 159-165, 1999).

- 10 The observation of elevated expression of Tie receptors in the endothelium of metastatic melanomas (Kaipainen et al., Cancer Res. 54, 6571-6577, 1994), in breast carcinomas (Salvén et al., Br. J. Cancer 74, 69-72, 1996), and in tumor xenografts grown in the presence of dominant-negative VEGF receptors (Millauer et al., Cancer Res. 56, 1615-1620, 1996), as well as elevated expression of Flt4 receptors in the endothelium of lymphatic vessels surrounding lymphomas and breast carcinomas (Jussila et al., Cancer Res. 58, 1599-1604, 1998), and of VEGF-C in various human tumor samples (Salvén et al., Am. J. Pathol. 153, 103-108, 1998), suggested these endothelium-specific growth factors and receptors as candidate alternative pathways driving tumor neovascularization. The high
- 15 upregulation of Ang2 expression already in early tumors has been interpreted in terms of a host defense mechanism against initial cooption of existing blood vessels by the developing tumor. In the absence of VEGF, the coopted vessels undergo regression leading to necrosis within the center of the tumor. Contrarily, hypoxic upregulation of VEGF expression in cooperation with elevated Ang2
- 20 expression rescues and supports tumor vascularization and tumor growth at the tumor margin (Holash et al., Science 284, 1994-1998, 1999; Holash et al., Oncogene 18, 5356-5362, 1999).

- 25 Interference with Tie2 receptor function by means of Angiopoietin-neutralizing Tie2 variants consisting of the extracellular ligand-binding domain has been shown to result in inhibition of growth and vascularization of experimental tumors (Lin et al., J. Clin. Invest. 103, 159-165, 1999; Lin et al. Proc. Natl. Acad. Sci. USA 95, 8829-8834, 1998; Siemeister et al., Cancer Res. 59, 3185-3191, 1999). Comparing the effects of interference with the endothelium-specific receptor

tyrosine kinase pathways by means of paracrine expression of the respective extracellular receptor domains on the same cellular background demonstrated inhibition of tumor growth upon blockade of the VEGF receptor system and of the Tie2 receptor system, respectively (Siemeister et al., Cancer Res. 59, 3185-3191, 5 1999).

It is known that the inhibition of the VEGF/VEGR receptor system by various methods resulted only in slowing down growth of most experimental tumors (Millauer et al., Nature 367, 576-579, 1994; Kim et al., Nature 362, 841-844, 1993; Millauer et al., Cancer Res. 56, 1615-1620, 1996; Goldman et al., Proc. Natl.

10 Acad. Sci. USA 95, 8795-8800, 1998; Fong et al., Cancer Res. 59, 99-106, 1999; Wedge et al., Cancer Res. 60, 970-975, 2000; Wood et al. Cancer Res. 60, 2178-2189, 2000; Siemeister et al., Cancer Res. 59, 3185-3191, 1999). Even by escalation of therapeutic doses a plateau level of therapeutic efficacy was achieved (Kim et al., Nature 362, 841-844, 1993; Wood et al. Cancer Res. 60, 15 2178-2189, 2000). Similar results were observed upon interference with the Angiopoietin/Tie2 receptor system (Lin et al., J. Clin. Invest. 103, 159-165, 1999; Lin et al., Proc. Natl. Acad. Sci. USA 95, 8829-8834, 1998; Siemeister et al., Cancer Res. 59, 3185-3191, 1999).

20 However, there is a high demand for methods that enhance the therapeutic efficacy of anti-angiogenous compounds.

Searching for methods that enhance the therapeutic efficacy of anti-angiogenic compounds, superior anti-tumor effects were observed unexpectedly upon 25 combination of inhibition of VEGF/VEGF receptor systems and interference with biological function of Angiopoietin/Tie receptor systems. The mode of action underlying the superior effects observed may be that interference biological function of Angiopoietin/Tie receptor systems destabilizes endothelial cell-periendothelial cell interaction of existing mature tumor vessels and thereby 30 sensitizes the endothelium to compounds directed against VEGF/VEGF receptor systems.

Based on this unexpected finding the present invention provides the combination of functional interference with VEGF/VEGF receptor systems and with

Angiopoietin/Tie receptor systems for inhibition of vascularization and of tumor growth.

The pharmaceutical composition consists of two components: compound I inhibits the biological activity of one or several of the VEGF/VEGF receptor systems or

- 5 consists of cytotoxic agents which are targeted to the endothelium via recognition of VEGF/VEGF receptor systems. Compound II interferes with the biological function of one or several of Angiopoietin/Tie receptor systems or consists of cytotoxic agents which are targeted to the endothelium via recognition of Angiopoietin/Tie receptor systems. Alternatively, compound I inhibits the biological
10 activity of one or several of the VEGF/VEGF receptor systems or of the Angiopoietin/Tie receptor systems and compound II consists of cytotoxic agents which are targeted to the endothelium via recognition of one or several of the VEGF/VEGF receptor systems or of the Angiopoietin/Tie receptor systems.
Targeting or modulation of the biological activities of VEGF/VEGF receptor
15 systems and of Angiopietin/Tie receptor systems can be performed by

- (a) compounds which inhibit receptor tyrosine kinase activity,
- (b) compounds which inhibit ligand binding to receptors,
- (c) compounds which inhibit activation of intracellular signal pathways of the

- 20 receptors,
(d) compounds which inhibit or activate expression of a ligand or of a receptor of the VEGF or Tie receptor system,

- (e) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which target cytotoxic agents
25 or coagulation-inducing agents to the endothelium via recognition of VEGF/VEGF receptor or Angiopoietin/Tie receptor systems,

- (f) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which are targeted to the
30 endothelium and induce necrosis or apoptosis.

A compound comprised by compositions of the present invention can be a small molecular weight substance, an oligonucleotide, an oligopeptide, a recombinant protein, an antibody, or conjugates or fusionproteins thereof. An example of an inhibitor is a small molecular weight molecule which inactivates a receptor tyrosine

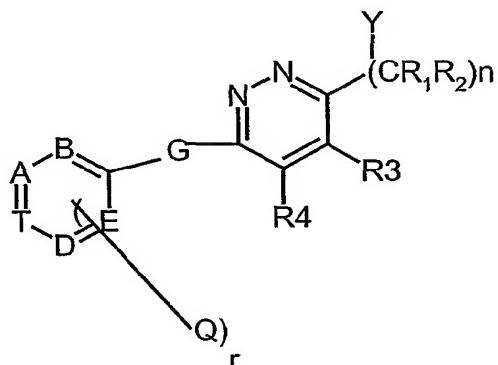
kinase by binding to and occupying the catalytic site such that the biological activity of the receptor is decreased. Kinase inhibitors are known in the art (Sugen: SU5416, SU6668; Fong et al. (1999), Cancer Res. 59, 99-106; Vajkoczy et al., Proc. Am. Assoc. Cancer Res. San Francisco (2000), Abstract ID 3612; Zeneca:

- 5 ZD4190, ZD6474; Wedge et al. (2000), Cancer Res. 60, 970-975; Parke-Davis PD0173073, PD0173074; Johnson et al., Proc. Am. Assoc. Cancer Res., San Francisco (2000), Abstract ID 3614; Dimitroff et al. (1999), Invest. New Drugs 17, 121-135). An example of an antagonist is a recombinant protein or an antibody which binds to a ligand such that activation of the receptor by the ligand is
10 prevented. Another example of an antagonist is an antibody which binds to the receptor such that activation of the receptor is prevented. An example of an expression modulator is an antisense RNA or ribozyme which controls expression of a ligand or a receptor. An example of a targeted cytotoxic agent is a fusion protein of a ligand with a bacterial or plant toxin such as Pseudomonas exotoxin
15 A, Diphtheria toxin, or Ricin A. An example of a targeted coagulation-inducing agent is a conjugate of a single chain antibody and tissue factor. Ligand-binding inhibitors such as neutralizing antibodies which are known in the art are described by Genentech (rhuMAbVEGF) and by Presta et al. (1997), Cancer Res. 57, 4593-4599. Ligand-binding receptor domaines are described by Kendall & Thomas
20 (1993), Proc. Natl. Acad. Sci., U.S.A.90, 10705-10709; by Goldman et al. (1998) Proc. Natl. Acad. Sci., U.S.A.95, 8795-8800 and by Lin et al. (1997), J. Clin. Invest. 100, 2072-2078. Further, dominant negative receptors have been described by Millauer et al. (1994), Nature 367, 567-579.
Receptor blocking antibodies have been described by Imclone (c-p1C11, US
25 5,874,542). Further known are antagonistic ligand mutants (Siemeister et al. (1998), Proc. Natl. Acad. Sci., U.S.A.95, 4625-4629). High affinity ligand- or receptor binding oligo nucleotides habe been described by NeXstar (NX-244) and Drolet et al. (1996), Nat. Biotech 14, 1021-1025. Further, small molecules and peptides have been described.
- 30 Expression regulators have been described as anti-sense oligo nucleotides and as ribozymes (RPI, Angiozyme™, see RPI Homepage).

Examples for delivery-/Targeting-Systems have been described as ligand/antibody-toxin-fusion-proteins or conjugates (Arora et al. (1999), Cancer Res. 59, 183-188 and Olson et al. (1997), Int. J. Cancer 73, 865-870), as endothel cell targeting of liposomes (Spragg et al. (1997), Prog. Natl. Acad. Sci., U.S.A94, 8795-8800, and as endothel cell targeting plus coagulation-induction (Ran et al., (1998), Cancer Res. 58, 4646-4653).

- 10 Small molecules which inhibit the receptor tyrosine kinase activity are for example molecules of general formula I

15



20

I,

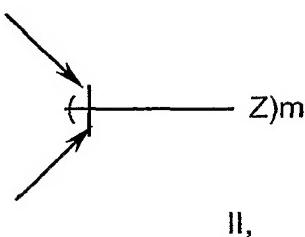
in which

r has the meaning of 0 to 2,
n has the meaning of 0 to 2;

25

R₃ und R₄ a) each independently from each other have the meaning of lower alkyl,

b) together form a bridge of general partial formula II;

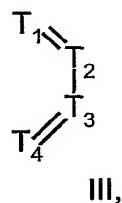


5

m wherein the binding is via the two terminal C- atoms, and has the meaning of 0 to 4; or

c) together form a bridge of partial formula III

10



15

wherein one or two of the ring members T₁, T₂, T₃, T₄ has the meaning of nitrogen, and each others have the meaning of CH, and the bining is via the atoms T₁ and T₄;

G

20

has the meaning of C₁ - C₆ - alkyl, C₂ - C₆ - alkylene or

C₂ - C₆ - alkenylene; or C₂ - C₆ - alkylene or C₃ - C₆ - alkenylene, which are substituted with acyloxy or hydroxy; -CH₂-O-, -CH₂-S-, -CH₂-NH-, -CH₂-O-CH₂-, -CH₂-S-CH₂-, -CH₂-NH-CH₂, oxa (-O-), thia (-S-) or imino (-NH-),

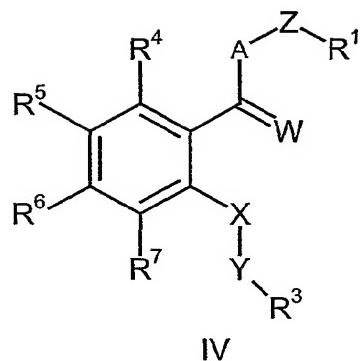
A, B, D, E and T

25

independently from each other have the meaning of N or CH, with the provisio that not more than three of these Substituents have the meaning of N,

- Q has the meaning of lower alkyl, lower alkyloxy or halogene,
R₁ and R₂ independently from each other have the meaning of H or
lower alkyl,
X has the meaning of imino, oxa or thia;
5 Y has the meaning of hydrogene, unsubstituted or substituted
aryl, heteroaryl, or unsubstituted or substituted cycloalkyl; and
Z has the meaning of amino, mono- or disubstituted amino,
halogen, alkyl, substituted alkyl, hydroxy, etherificated or
esterificated hydroxy, nitro, cyano, carboxy, esterificated
carboxy, alkanoyl, carbamoyl, N-mono- or N, N- disubstituted
10 carbamoyl, amidino, guanidino, mercapto, sulfo, phenylthio,
phenyl-lower-alkyl-thio, alkyl-phenyl-thio, phenylsulfinyl,
phenyl-lower-alkyl-sulfinyl, alkylphenylsulfinyl, phenylsulfonyl,
phenyl-lower-alkan-sulfonyl, or alkylphenylsulfonyl, whereas, if
15 more than one rest Z is present ($m \geq 2$), the substituents Z are
equal or different from each other, and wherein the bonds
marked with an arrow are single or double bonds; or an N-
oxide of said compound, wherein one ore more N-atoms carry
an oxygene atom, or a salt thereof.
- 20 A preferred salt is the salt of an organic acid, especially a succinate.
- These compounds can preferentially be used as compound I or II in the inventive
pharmaceutical composition.
- 25 Compounds which stop a tyrosin phosphorylation, or the persistent angiogenesis,
respectively, which results in a prevention of tumor growth and tumor spread, are
for example
anthranyl acid derivatives of general formula IV
- 30

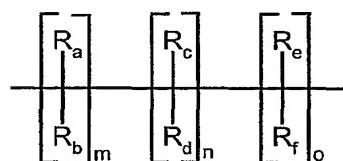
11



in which

- A has the meaning of group =NR²,
- 5 W has the meaning of oxygen, sulfur, two hydrogen atoms or the group =NR⁸,
- Z has the meaning of the group =NR¹⁰ or =N-, -N(R¹⁰)-
(CH₂)_q-, branched or unbranched C₁₋₆-Alkyl or is the group

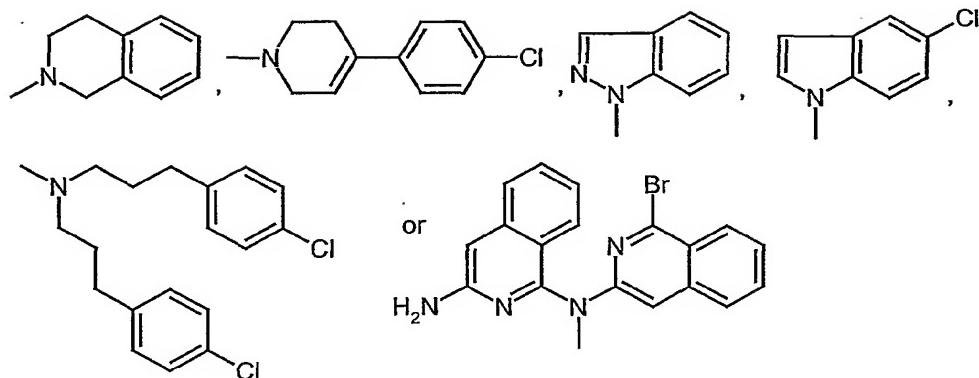
10



15

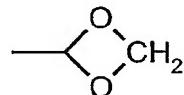
or A, Z and R¹ together form the group

20



| | | |
|----|---|---|
| | m, n and o | has the meaning of 0 – 3, |
| | q | has the meaning of 1 – 6, |
| 5 | R _a , R _b , R _c , R _d , R _e , R _f | independently from each other have the meaning of hydrogen, C ₁₋₄ alkyl or the group =NR ¹⁰ , and/or R _a and/or R _b together with R _c and/or R _d or R _c together with R _e and/or R _f form a bound, or up to two of the groups R _a -R _f form a bridge with each up to 3 C-atoms with R ¹ or R ² , |
| 10 | X | has the meaning of group =NR ⁹ or =N-, |
| | Y | has the meaning of group -(CH ₂) _p , |
| | p | has the meaning of integer 1-4, |
| | R ¹ | has the meaning of unsubstituted or optionally substituted with one or more of halogene, C ₁₋₆ -alkyl, or C ₁₋₆ -alkyl or C ₁₋₆ -alkoxy, which is optionally substituted by one or more of halogen, or is unsubstituted or substituted aryl or heteroaryl, |
| 15 | R ² | has the meaning of hydrogen or C ₁₋₆ -alkyl, or form a bridge with up to 3 ring atoms with R _a -R _f together with Z or R ₁ , |
| 20 | R ³ | has the meaning of monocyclic or bicyclic aryl or heteroaryl which is unsubstituted or optionally substituted with one or more of halogen, C ₁₋₆ -alkyl, C ₁₋₆ -alkoxy or hydroxy, |
| 25 | R ⁴ , R ⁵ , R ⁶ and R ⁷ | independently from each other have the meaning of hydrogen, halogen or C ₁₋₆ -alkoxy, C ₁₋₆ -alkyl or |

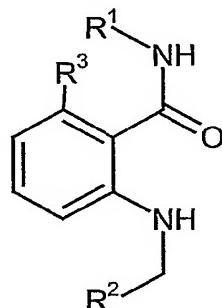
C_{1-6} -carboxyalkyl, which are unsubstituted or optionally substituted with one or more of halogen, or R^5 and R^6 together form the group



- 5 R^8 , R^9 and R^{10} independently from each other have the meaning of hydrogen or C_{1-6} -alkyl,
as well as their isomers and salts.

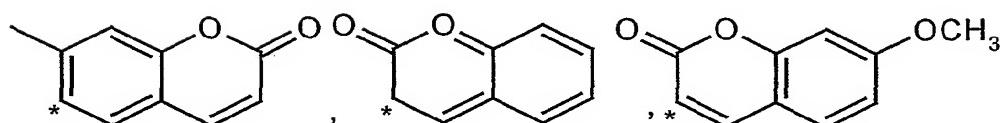
These compounds can also preferentially be used as compound I or II in the
10 inventive pharmaceutical composition.

More preferentially compounds of general formula V

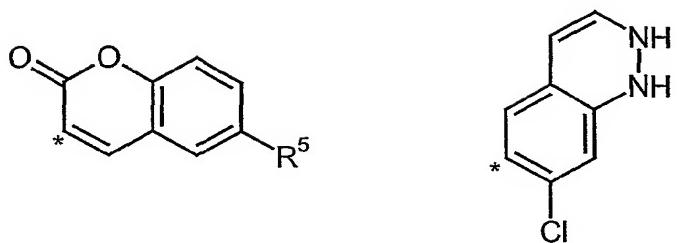


- 15 V,
in which
 R^1 has the meaning of group

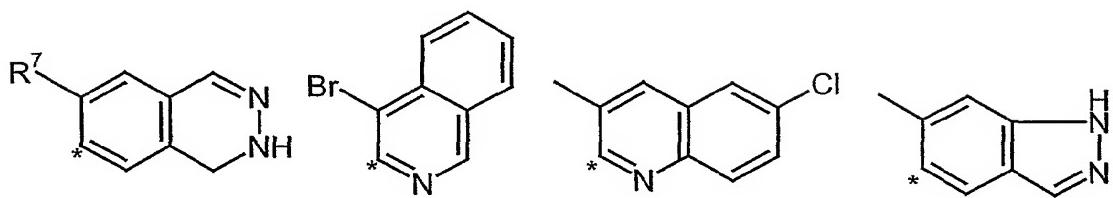
20



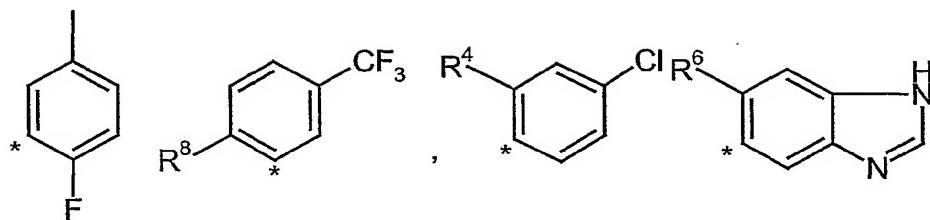
14



in which R⁵ is chloro, bromo or the group -OCH₃,



in which R⁷ is -CH₃ or chloro,



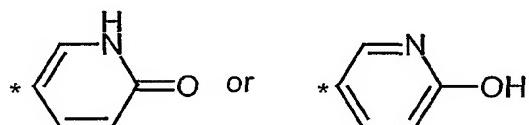
in which R⁸ is -CH₃, fluoro,
chloro or -CF₃

in which R⁴ is fluoro,
chloro, bromo, -CF₃,
-N≡C, -CH₃, -OCF₃ or
-CH₂OH

in which R⁶ is
-CH₃ or chloro

5

R² has the meaning of pyridyl or the group



10

and

R³ has the meaning of hydrogen or fluoro, as well as their isomers and salts can be used as compound I or II in the inventive pharmaceutical composition.

These compounds have the same properties as already mentioned above under 15 compound IV and can be used for the treatment of angiogeneous diseases.

Compositions comprise compounds of general formulars I, IV and V, alone or in combination.

The above mentioned compounds are also claimed matter within the inventive combinations.

20

A further example for ligand binding inhibitors are peptides and DNA sequences coding for such peptides, which are used for the treatment of angiogeneous diseases. Such peptides and DNA sequences are disclosed in Seq. ID No. 1 to 59 of the sequence protocoll. It has been shown that Seq. ID Nos. 34 and 34a are of 25 main interest.

Claimed matter of the instant invention are therefor pharmaceutical compositions

- a) comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems,
- 5 b) comprising one or several agents as compound I which are targeted to the endothelium via one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems,
- 10 c) comprising one or several agents as compound I which modulates the biological function of one or several of the VEGF/VEGF receptor systems or of one or several of the Angiopoietin/ Tie receptor systems and comprising one or several agents as compound II which are targeted to the endothelium,
- 15 d) comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems,
- 20 e) comprising one or several agents as compound I which are targeted to the endothelium via one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems,
- 25 f) comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the VEGF/VEGF receptor systems,
- 30 g) comprising one or several agents as compound I which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems and

h) comprising one or several agents which interfere with both the function of one or several of the VEGF/VEGF receptor systems and the function of one or several of the Angiopoietin/Tie receptor systems.

5

For a sequential therapeutical application the inventive pharmaceutical compositions can be applied simultaneously or separately .

The inventive compositions comprise as compound I or as compound II at least

10 one of

- a) compounds which inhibit receptor tyrosine kinase activity,
- b) compounds which inhibit ligand binding to receptors,
- c) compounds which inhibit activation of intracellular signal pathways of the receptors,
- 15 d) compounds which inhibit or activate expression of a ligand or of a receptor of the VEGF or Tie receptor system,
- e) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which target cytotoxic agents or coagulation-inducing agents to the endothelium via recognition of
- 20 VEGF/VEGF receptor or Angiopoietin/Tie receptor systems,
- f) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which are targeted to the endothelium and induce necrosis or apoptosis.

These compositions are also claimed matter of the present invention.

25

Also claimed matter of the present invention are pharmaceutical compositions which comprise as compound I and/ or II at least one of Seq. ID Nos. 1-59.

Of most value are pharmaceutical compositions, which comprise as compound I and/ or II Seq. ID Nos. 34a und pharmaceutical compositions according to claims 30 which comprise as compound I and/ or II at least one of sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate.

Further preferred matter of the present invention are pharmaceutical compositions, which comprise as compound I and/ or II at least one small molecule of general formula I, general formula IV and/ or general formula V.

- 5 The most preferred compound which can be used as compound I or II in the inventive composition is (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate.
Therefore, claimed matter of the present invention are also pharmaceutical compositions, which comprise as compound I (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate, sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate, and as compound II (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate, Tie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate, with the proviso that compound I is not identically to compound II, and most preferred
10 pharmaceutical compositions, which comprise as compound I (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate and as compound II sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate; pharmaceutical compositions, which comprise as compound I mAB 4301-42-35 and as compound II sTie2, and/ or scFv-tTF conjugate; pharmaceutical
15 compositions, which comprise as compound I scFv-tTF conjugate and as compound II sTie2 and/ or mAB 4301-42-35; pharmaceutical compositions, which comprise as compound I L19 scFv-tTF conjugate and as compound II sTie2.

- The small molecule compounds, proteins and DNA's expressing proteins, as
20 mentioned above can be used as medicament alone, or in form of formulations for the treatment of tumors, cancers, psoriasis, arthritis, such as rheumatoide arthritis, hemangioma, angiofibroma, eye diseases, such as diabetic retinopathy, neovascular glaukoma, kidney diseases, such as glomerulonephritis, diabetic nephropathy, malignant nephrosclerose, thrombotic microangiopathic syndrome,
25 transplantation rejections and glomerulopathy, fibrotic diseases, such as cirrhotic liver, mesangial cell proliferative diseases, arteriosclerosis and damage of nerve tissues.

The treatment of the damaged nerve tissues with the inventive combination hinders the rapid formation of scars at the damaged position. Thus, there is no

scar formation before the axons communicate with each other. Therefore a reconstruction of the nerve bindings is much more easier.

Further, the inventive combinations can be used for suppression of the ascites

5 formation in patients. It is also possible to suppress VEGF oedemas.

For the use of the inventive combinations as medicament the compounds will be formulated as pharmaceutical composition. Said formulation comprises beside the active compound or compounds acceptable pharmaceutically, organically or inorganically inert carriers, such as water, gelatine, gum arabic, lactose, starch,

10 magnesium stearate, talcum, plant oils, polyalkylene glycols, etc. Said

pharmaceutical preparations can be applied in solid form, such as tablets, pills, suppositories, capsules, or can be applied in fluid form, such as solutions, suspensions or emulsions.

If necessary, the compositions additionally contain additives, such as

15 preservatives, stabilizer, detergents or emulgators, salts for alteration of the osmotic pressure and/ or buffer.

These uses are also claimed matter of the instant invention, as well as the formulations of the active compounds

20 For parenteral application especially injectable solutions or suspensions are suitable, especially hydrous solutions of the active compound in polyhydroxyethoxylated castor-oil are suitable.

As carrier also additives can be used, such as salts of the gallic acid or animal or plant phospholipids, as well as mixtures thereof, and liposomes or ingredients 25 thereof.

For oral application especially suitable are tablets, pills or capsules with talcum and/ or hydrocarbon carriers or binders, such as lactose, maize or potato starch.

The oral application can also be in form of a liquid, such as juice, which optionally contains a sweetener.

30 The dosis of the active compound differs depending on the application of the compound, age and weight of the patient, as well as the form and the progress of the disease.

The daily dosage of the active compound is 0,5-1000 mg, especially 50-200 mg.

The dosis can be applied as single dose or as two or more daily dosis.

These formulations and application forms are also part of the instant invention.

Combined functional interference with VEGF/VEGF receptor systems and with

- 5 Angiopoietin/Tie receptor systems can be performed simultaneously, or in sequential order such that the biological response to interference with one ligand/receptor system overlaps with the biological response to interference with a second ligand/receptor system. Alternatively, combined functional interference with VEGF/VEGF receptor systems or with Angiopoietin/Tie receptor systems and
- 10 targeting of cytotoxic agents via VEGF/VEGF receptor systems or via Angiopoietin/Tie receptor systems can be performed simultaneously, or in sequential order such that the biological response to functional interference with a ligand/receptor system overlaps in time with targeting of cytotoxic agents.
- 15 The invention is also directed to a substance which functional interferes with both VEGF/VEGF receptor systems and Angiopoietin/Tie receptor systems, or which are targeted via both VEGF/VEGF receptor systems and Angiopoietin/Tie receptor systems.
- 20 VEGF/VEGF receptor systems include the ligands VEGF-A, VEGF-B, VEGF-C, VEGF-D, PIGF, and the receptor tyrosine kinases VEGF-R1 (Flt1), VEGF-R2 (KDR/Flik1), VEGF-R3 (Flt4), and their co-receptors (i.e. neuropilin-1). Angiopoietin/Tie receptor systems include Ang1, Ang2, Ang3/Ang4, and angiopoietin related polypeptides which bind to Tie1 or to Tie2, and the receptor
- 25 tyrosine kinases Tie1 and Tie2.

Pharmaceutical compositions of the present invention can be used for medicinal purposes. Such diseases are, for example, cancer, cancer metastasis, angiogenesis including retinopathy and psoriasis. Pharmaceutical compositions of

- 30 the present invention can be applied orally, parenterally, or via gene therapeutic methods.

Therefor the present invention also concerns the use of pharmaceutical compositions for the production of a medicament for the treatment of tumors,

cancers, psoriasis, arthritis, such as rheumatoide arthritis, hemangioma, angiofibroma, eye diseases, such as diabetic retinopathy, neovascular glaucoma, kidney diseases, such as glomerulonephritis, diabetic nephropathie, malignant nephrosclerosis, thrombic microangiopathic syndrome, transplantation rejections
5 and glomerulopathy, fibrotic diseases, such as cirrhotic liver, mesangial cell proliferative diseases, arteriosclerosis, damage of nerve tissues, suppression of the ascites formation in patients and suppression of VEGF oedemas.

The following examples demonstrate the feasibility of the disclosed invention, without restricting the invention to the disclosed examples.

5 **Example 1**

Superior effect on inhibition of tumor growth via combination of inhibition of the VEGF A/VEGF receptor system together with functional interference with the Angiopoietin/Tie2 receptor system over separate modes of intervention was demonstrated in an A375v human melanoma xenograft model.

10

Human melanoma cell line A375v was stably transfected to overexpress the extracellular ligand-neutralizing domain of human Tie2 receptor tyrosine kinase (sTie2; compound II) (Siemeister et al., Cancer Res. 59, 3185-3191, 1999). For control, A375v cells were stably transfected with the empty expression vector

15

(A375v/pCEP). Swiss *nu/nu* mice were s.c. injected with 1×10^6 transfected A375v/sTie2 or A375v/pCEP tumor cells, respectively. Animals receiving compound I were treated for up to 38 days with daily oral doses of 50 mg/kg of the

VEGF receptor tyrosine kinase inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (Wood et al., Cancer Res. 60,

20

2178-2189, 2000). Various modes of treatment are described in Table 1. Tumor growth was determined by caliper measurement of the largest diameter and its perpendicular.

Table 1

| | mode of treatment | |
|-------------------------|--|------------------------|
| treatment group | (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (compound I) | sTie2 (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/sTie2 | - | + |
| Group 4: A375v/sTie2 | + | + |

- 5 Tumors derived from A375v/pCEP control cells reached a size of approx. 250 mm² (mean area) within 24 days (Figure 1) without treatment (group 1). Separate treatment with the VEGF receptor inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (compound I, treatment group 2) or separate interference with Angiopoietin/Tie2 receptor system by means of expression of sTie2 (compound II, treatment group 3) delayed growth of tumors to a size of approx. 250 mm² to 31 days, respectively. Combination of interference with the Angiopoietin/Tie2 system by means of expression of sTie2 and of interference with the VEGF/VEGF receptor system by means of the kinase inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (compound I + compound II, treatment group 4) delayed growth of the tumors to a size of approx. 250 mm² to 38 days.
- This result clearly demonstrates the superior effect of a combination of interference with the VEGF-A/VEGF receptor system and the Angiopoietin/Tie2 receptor system over separate modes of intervention.

Example 2

Combination of functional interference with the Angiopoietin/Tie2 receptor system and neutralization of VEGF-A is superior to separate modes of intervention in
 5 inhibition of tumor growth.

Tumors derived from A375v/sTie2 cells and from A375v/pCEP cells were induced in nude mice as described in example 1. Animals receiving compound I were treated twice weekly over a period of time of 4 weeks with intraperitoneal doses of
 10 200 µg of the VEGF-A-neutralizing monoclonal antibody (mAb) 4301-42-35 (Schlaeppi et al., J. Cancer Res. Clin. Oncol. 125, 336-342, 1999). Various modes of treatment are described in Table 2. Animals were sacrificed for ethical reasons when tumors of group 1 exceeded a volume of approx. 1000 mm³. Tumor growth was determined by caliper measurement of the largest diameter and its
 15 perpendicular.

Table 2

| treatment group | mode of treatment | |
|-------------------------|--------------------------------|------------------------|
| | mAb 4301-42-35 (compound I) | sTie2 (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/sTie2 | - | + |
| Group 4: A375v/sTie2 | + | + |

Tumors derived from A375v/pCEP control cells reached a size of approx. 1000
 20 mm³ within 28 days (Figure 2) without treatment (group 1). Tumors treated with the VEGF-A-neutralizing mAb 4301-42-35 (compound I, treatment group 2) grew to a volume of approx. 450 mm³ within 28 days. Interference with

Angiopoietin/Tie2 receptor system by means of expression of sTie2 (compound II, treatment group 3) reduced growth of tumors within 28 day to a volume of approx. 600 mm², respectively. Combination of interference with the Angiopoietin/Tie2

system by means of expression of sTie2 and neutralizing of VEGF-A by means of

- 5 the mAb 4301-42-35 (compound I + compound II, treatment group 4) resulted in a inhibition of tumor growth to a volume of approx. 250 mm³ within 28 days.

The superior effect of a combination of neutralization of VEGF-A and functional interference with the Angiopoietin/Tie2 receptor system over separate modes of

- 10 intervention is clearly shown.

Example 3

Combination of functional interference with the Angiopoietin/Tie2 receptor system and targeting of a coagulation-inducing protein via the VEGF/VEGF receptor system is superior to separate modes of intervention in inhibition of tumor growth.

5

Tumors derived from A375v/sTie2 cells and from A375v/pCEP cells were induced in nude mice as described in example 1. A single chain antibody (scFv) specifically recognizing the human VEGF-A/VEGF receptor I complex (WO 99/19361) was expressed in E. coli and conjugated to coagulation-inducing

10 recombinant human truncated tissue factor (tTF) by methods described by Ran et al. (Cancer Res. 58, 4646-4653, 1998). When tumors reached a size of approx. 200 mm³ animals receiving compound I were treated on day 0 and on day 4 with intravenous doses of 20 µg of the scFv-tTF conjugate. Various modes of treatment are described in Table 3. Animals were sacrificed for ethical reasons
 15 when tumors of group 1 exceeded a volume of approx. 1000 mm³. Tumor growth was determined by caliper measurement of the largest diameter and its perpendicular.

Table 3

| treatment group | mode of treatment | |
|-------------------------|------------------------------------|------------------------|
| | scFv-tTF conjugate (compound I) | sTie2 (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/sTie2 | - | + |
| Group 4: A375v/sTie2 | + | + |

Tumors derived from A375v/pCEP control cells reached a size of approx. 1000 mm³ within 28 days (Figure 3) without treatment (group 1). Tumors treated with the coagulation-inducting tTF targeted to the VEGF-A/VEGF receptor I complex via the scFv-tTF conjugate (compound I, treatment group 2) grew to a volume of
5 approx. 500 mm³ within 28 days. Interference with Angiopoietin/Tie2 receptor system by means of expression of sTie2 (compound II, treatment group 3) reduced growth of tumors within 28 day to a volume of approx. 600 mm², respectively. Combination of interference with the Angiopoietin/Tie2 system by means of expression of sTie2 and of targeting the VEGF receptor complex
10 (compound I + compound II, treatment group 4) resulted in a inhibition of tumor growth to a volume of approx. 300 mm³ within 28 days.
The superior effect of a combination of targeting of the coagulation-inducting tTF to the VEGF-A/VEGF receptor I complex and functional interference with the Angiopoietin/Tie2 receptor system over separate modes of intervention is clearly
15 shown. Similar effects can be expected upon targeting of cytotoxic agents to VEGF/VEGF receptor systems.

Example 4

Combination of functional interference with the VEGF/VEGF receptor system and targeting of a coagulation-inducing protein via the VEGF/VEGF receptor system is

- 5 superior to separate modes of intervention in inhibition of tumor growth.

Tumors derived from A375v/pCEP cells were induced in nude mice as described in example 1. Animals receiving compound I were treated for up to 28 days with daily oral doses of 50 mg/kg of the VEGF receptor tyrosine kinase inhibitor (4-

- 10 Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (Wood et al., Cancer Res. 60, 2178-2189, 2000). Compound II consists of a single chain antibody (scFv) specifically recognizing the human VEGF-A/VEGF receptor I complex (WO 99/19361) which was expressed in E. coli and conjugated to coagulation-inducing recombinant human truncated tissue factor (tTF) by methods
15 described by Ran et al. (Cancer Res. 58, 4646-4653, 1998). When tumors reached a size of approx. 200 mm³ animals receiving compound II were treated on day 0 and on day 4 with intravenous doses of 20 µg of the scFv-tTF conjugate. Various modes of treatment are described in Table 4. Animals were sacrificed for ethical reasons when tumors of group 1 exceeded a volume of approx. 1000 mm³. Tumor
20 growth was determined by caliper measurement of the largest diameter and its perpendicular.

Table 4

| | mode of treatment | |
|------------------------|---|-------------------------------------|
| treatment group | (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthal-azin-1-yl]ammonium hydrogen succinate (compound I) | scFv-tTF conjugate (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/pCEP | - | + |
| Group 4: A375v/pCEP | + | + |

- 5 Tumors derived from A375v/pCEP control cells reached a size of approx. 1000 mm³ within 28 days (Figure 4) without treatment (group 1). Separate treatment with the VEGF receptor inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (compound I, treatment group 2) resulted in a reduction of the tumor volumes to approx. 550 mm³. Tumors treated with the
- 10 coagulation-inducting tTF targeted to the VEGF-A/VEGF receptor I complex via the scFv-tTF conjugate (compound II, treatment group 3) grew to a volume of approx. 500 mm³ within 28 days. Combination of inhibition of VEGF receptor tyrosine kinase by means of (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate and of targeting the VEGF receptor complex
- 15 (compound I + compound II, treatment group 4) resulted in a inhibition of tumor growth to a volume of approx. 400 mm³ within 28 days.

The superior effect of a combination of targeting of the coagulation-inducting tTF to the VEGF-A/VEGF receptor I complex and functional interference with the

20 VEGF/VEGF receptor system over separate modes of intervention is clearly

shown. Similar effects can be expected upon targeting of cytotoxic agents to Angiopoietin/Tie receptor systems.

Example 5

Combination of functional interference with the Angiopoietin/Tie2 receptor system and endothelium-specific targeting of a coagulation-inducing protein is superior to
 5 separate modes of intervention in inhibition of tumor growth.

Tumors derived from A375v/sTie2 cells and from A375v/pCEP cells were induced in nude mice as described in example 1. A fusion protein (L19 scFv-tTF) consisting of L19 single chain antibody specifically recognizing the oncofoetal ED-
 10 B domain of fibronectin and the extracellular domain of tissue factor was expressed in E. coli as described by Nilsson et al. (Nat. Med., in press). Further, L19 scFv-tTF data have been represented by D. Neri and F. Nilsson (Meeting "Advances in the application of monoclonal antibodies in clinical oncology", Samos, Greece, 31. May-2. June 2000). When tumors reached a size of approx.
 15 200 mm³ animals receiving compound I were treated with a single intravenous dose of 20 µg of L19 scFv-tTF in 200 µl saline. Various modes of treatment are described in Table 5. Animals were sacrificed for ethical reasons when tumors of group 1 exceeded a volume of approx. 1000 mm³. Tumor growth was determined by caliper measurement of the largest diameter and its perpendicular.

20

Table 5

| treatment group | mode of treatment | |
|-------------------------|------------------------------|------------------------|
| | L19 scFv-tTF (compound I) | sTie2 (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/sTie2 | - | + |
| Group 4: A375v/sTie2 | + | + |

Tumors derived from A375v/pCEP control cells reached a size of approx. 1000 mm³ within 28 days (Figure 5) without treatment (group 1). Tumors treated with the coagulation-inducting L19 scFv-tTF (compound I, treatment group 2) grew to a 5 volume of approx. 450 mm³ within 28 days. Interference with Angiopoietin/Tie2 receptor system by means of expression of sTie2 (compound II, treatment group 3) reduced growth of tumors within 28 day to a volume of approx. 600 mm², respectively. Combination of interference with the Angiopoietin/Tie2 system by means of expression of sTie2 and of targeting the endothelium with L19 scFv-tTF 10 (compound I + compound II, treatment group 4) resulted in a inhibition of tumor growth to a volume of approx. 250 mm³ within 28 days.

The superior effect of a combination of targeting of L19 scFv-tTF to the endothelium and functional interference with the Angiopoietin/Tie2 receptor system over separate modes of intervention is clearly shown.

Example 6

Combination of functional interference with the VEGF/VEGF receptor system and endothelium-specific targeting of a coagulation-inducing protein is superior to
5 separate modes of intervention in inhibition of tumor growth.

Tumors derived from A375v/pCEP cells were induced in nude mice as described in example 1. Animals receiving compound I were treated for up to 28 days with daily oral doses of 50 mg/kg of the VEGF receptor tyrosine kinase inhibitor (4-
10 Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (Wood et al., Cancer Res. 60, 2178-2189, 2000). Compound II consists of L19 scFv-tTF fusion protein as described in example 5. When tumors reached a size of approx. 200 mm³ animals receiving compound II were treated with a single intravenous dose of 20 µg of L19 scFv-tTF in 200 µl saline. Various modes of
15 treatment are described in Table 6. Animals were sacrificed for ethical reasons when tumors of group 1 exceeded a volume of approx. 1000 mm³. Tumor growth was determined by caliper measurement of the largest diameter and its perpendicular.

Table 6

| treatment group | mode of treatment | |
|------------------------|---|-------------------------------|
| | (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthal-azin-1-yl]ammonium hydrogen succinate (compound I) | L19 scFv-tTF (compound II) |
| Group 1: A375v/pCEP | - | - |
| Group 2: A375v/pCEP | + | - |
| Group 3: A375v/pCEP | - | + |
| Group 4: A375v/pCEP | + | + |

5

Tumors derived from A375v/pCEP control cells reached a size of approx. 1000 mm³ within 28 days (Figure 6) without treatment (group 1). Separate treatment with the VEGF receptor inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate (compound I, treatment group 2) resulted in a

10 reduction of the tumor volumes to approx. 550 mm³. Tumors treated with the coagulation-inducting L19 scFv-tTF targeted to the endothelium (compound II, treatment group 3) grew to a volume of approx. 450 mm³ within 28 days.

Combination of inhibition of VEGF receptor tyrosine kinase by means of (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate

15 and of targeting the VEGF receptor complex (compound I + compound II, treatment group 4) resulted in a inhibition of tumor growth to a volume of approx. 200 mm³ within 28 days.

The superior effect of a combination of targeting of L19 scFv-tTF to the endothelium and functional interference with the VEGF/VEGF receptor system over separate modes of intervention is clearly shown.

5

10

Description of the figures

Fig. 1 shows the superior effect of combination of interference with VEGF/VEGF receptor system by means of an specific tyrosine kinase inhibitor and with the

- 5 Angiopoietin/Tie2 receptor system by means of a soluble receptor domain on inhibition of tumor growth (treatment modes of groups 1-4 are given in Table 1).

The abbreviations have the following meaning:

| | | |
|-------------------|---|-------------------|
| mock, con. | = | treatment group 1 |
| mock+VEGF-A | = | treatment group 2 |
| 10 sTIE2-cl13 | = | treatment group 3 |
| sTIE2-cl13+VEGF-A | = | treatment group 4 |

Fig. 2 shows the superior effect on tumor growth inhibition of combination of

- 15 VEGF-neutralization and functional interference with Angiopoietin/Tie2 receptor system over separate modes of intervention (treatment modes of groups 1-4 are given in Table 2).

- 20 Fig. 3 shows the superior effect on tumor growth inhibition of combination of targeting of the coagulation-inducing tTF to the VEGF/VEGF receptor I complex via a scFv-tTF conjugate and functional interference with Angiopoietin/Tie2 receptor system over separate modes of intervention (treatment modes of groups 1-4 are given in Table 3).

25

Fig. 4 shows the superior effect on tumor growth inhibition of combination of targeting of the coagulation-inducing tTF to the VEGF/VEGF receptor I complex via a scFv-tTF conjugate and functional interference with VEGF/VEGF receptor system by means of the VEGF receptor tyrosine kinase inhibitor (4-

- 30 Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate over separate modes of intervention (treatment modes of groups 1-4 are given in Table 4).

Fig. 5 shows the superior effect on tumor growth inhibition of combination of targeting of the coagulation-inducing L19 scFv-tTF fusion protein to the endothelium and functional interference with Angiopoietin/Tie2 receptor system over separate modes of intervention (treatment modes of groups 1-4 are given in 5 Table 5).

Fig. 6 shows the superior effect on tumor growth inhibition of combination of targeting of the coagulation-inducing L19 scFv-tTF fusion protein to the endothelium and functional interference with VEGF/VEGF receptor system by means of the VEGF receptor tyrosine 10 kinase inhibitor (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate over separate modes of intervention (treatment modes of groups 1-4 are given in Table 6).

CLAIMS

1. Pharmaceutical compositions comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems.
5
2. Pharmaceutical compositions comprising one or several agents as compound I which are targeted to the endothelium via of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems.
10
3. Pharmaceutical compositions comprising one or several agents as compound I which modulates the biological function of one or several of the VEGF/VEGF receptor systems or of one or several of the Angiopoietin/ Tie receptor systems and comprising one or several agents as compound II which are targeted to the endothelium.
15
4. Pharmaceutical compositions comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems.
20
5. Pharmaceutical compositions comprising one or several agents as compound I which are targeted to the endothelium via one or several of the VEGF/VEGF receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems.
25
6. Pharmaceutical compositions comprising one or several agents as compound I which modulate the biological function of one or several of the VEGF/VEGF
30

receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the VEGF/VEGF receptor systems.

5 7. Pharmaceutical compositions comprising one or several agents as compound I which modulate the biological function of one or several of the Angiopoietin/Tie receptor systems, and comprising one or several agents as compound II which are targeted to the endothelium via one or several of the Angiopoietin/Tie receptor systems.

10

8. Pharmaceutical compositions comprising one or several agents which interfere with both the function of one or several of the VEGF/VEGF receptor systems and the function of one or several of the Angiopoietin/Tie receptor systems.

15 9. Pharmaceutical compositions according to claims 1-8 which are intended for simultaneous or separate sequential therapeutical application.

10. Pharmaceutical compositions according to claims 1-8 which comprise as compound I at least one of

- 20 a) compounds which inhibit receptor tyrosine kinase activity,
- b) compounds which inhibit ligand binding to receptors,
- c) compounds which inhibit activation of intracellular signal pathways of the receptors,
- d) compounds which inhibit or activate expression of a ligand or of a receptor of the VEGF or Tie receptor system,
- 25 e) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which target cytotoxic agents or coagulation-inducing agents to the endothelium via recognition of VEGF/VEGF receptor or Angiopoietin/Tie receptor systems,
- 30 f) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which are targeted to the endothelium and induce necrosis or apoptosis.

11. Pharmaceutical compositions according to claims 1-8 which comprise as compound II at least one of

- g) compounds which inhibit receptor tyrosine kinase activity,
- h) compounds which inhibit ligand binding to receptors,
- 5 i) compounds which inhibit activation of intracellular signal pathways of the receptors,
- j) compounds which inhibit or activate expression of a ligand or of a receptor of the VEGF or Tie receptor system,
- k) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which target cytotoxic agents or coagulation-inducing agents to the endothelium via recognition of VEGF/VEGF receptor or Angiopoietin/Tie receptor systems,
- 10 l) delivery systems, such as antibodies, ligands, high-affinity binding oligonucleotides or oligopeptides, or liposomes, which are targeted to the endothelium and induce necrosis or apoptosis.
- 15

12. Pharmaceutical compositions according to claims 1-11 which comprise as compound I and/ or II at least one of Seq. ID Nos. 1-59.

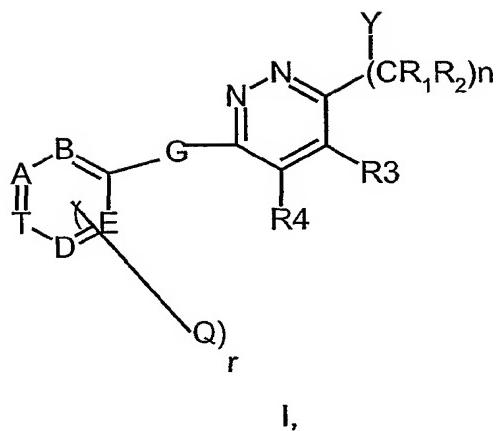
20

13. Pharmaceutical compositions according to claims 1-11 which comprise as compound I and/ or II Seq. ID Nos. 34a

14. Pharmaceutical compositions according to claims 1-11 which comprise as compound I and/ or II at least one of sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTFconjugate.

15. Pharmaceutical compositions according to claims 1-11 which comprise as compound I and/ or II at least one small molecule of genaral formula I

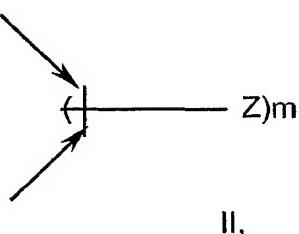
41



in which

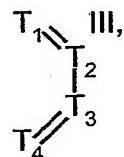
- 5 r has the meaning of 0 to 2,
 n has the meaning of 0 to 2;

- R₃ und R₄ a) each independently from each other have the
 meaning of lower alkyl,
 10 b) together form a bridge of general partial formula
 II,



- 15 wherein the binding is via the two terminal C- atoms,
 and
 m has the meaning of 0 to 4; or
 c) together form a bridge of partial formula III

20

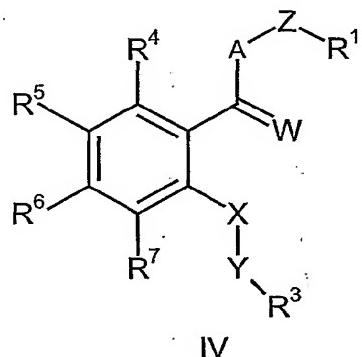


- wherein one or two of the ring members T_1, T_2, T_3, T_4 has the meaning of nitrogen, and each others have the meaning of CH, and the bining is via the atoms T_1 and T_4 ;
- 5 G has the meaning of $C_1 - C_6$ - alkyl, $C_2 - C_6$ – alkylene or $C_2 - C_6$ – alkenylene; or $C_2 - C_6$ - alkylene or $C_3 - C_6$ - alkenylene, which are substituted with acyloxy or hydroxy; $-CH_2-O-$, $-CH_2-S-$, $-CH_2-NH-$, $-CH_2-O-CH_2-$, $-CH_2-S-CH_2-$, $-CH_2-NH-CH_2$, oxa (-O-), thia (-S-) or imino (-NH-),
- 10 A, B, D, E and T independently from each other have the meaning of N or CH , with the provisio that not more than three of these Substituents have the meaning of N,
- 15 Q has the meaning of lower alkyl, lower alkyloxy or halogene,
- R₁ and R₂ independently from each other have the meaning of H or lower alkyl,
- 20 X has the meaning of imino, oxa or thia;
- Y has the meaning of hydrogene, unsubstituted or substituted aryl, heteroaryl, or unsubstituted or substituted cycloalkyl; and
- 25 Z has the meaning of amino, mono- or disubstituted amino, halogen, alkyl, substituted alkyl, hydroxy, etherificated or esterificated hydroxy, nitro, cyano, carboxy, esterificated carboxy, alkanoyl, carbamoyl, N-mono- or N, N- disubstituted carbamoyl, amidino, guanidino, mercapto, sulfo, phenylthio, phenyl-lower-alkyl-thio, alkyl-phenyl-thio, phenylsulfinyl, phenyl-lower-alkyl-sulfinyl, alkylphenylsulfinyl, phenylsulfonyl, phenyl-lower-alkan-sulfonyl, or alkylphenylsulfonyl, whereas, if more than one rest Z is present ($m \geq 2$), the substituents Z are equal or different from each other, and wherein the bonds marked with an arrow are single
- 30

or double bonds; or an N-oxide of said compound,
wherein one ore more N-atoms carry an oxygene atom,
or a salt thereof,

and/or a compound of genaral formula IV

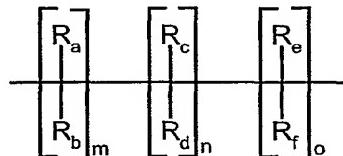
5



in which

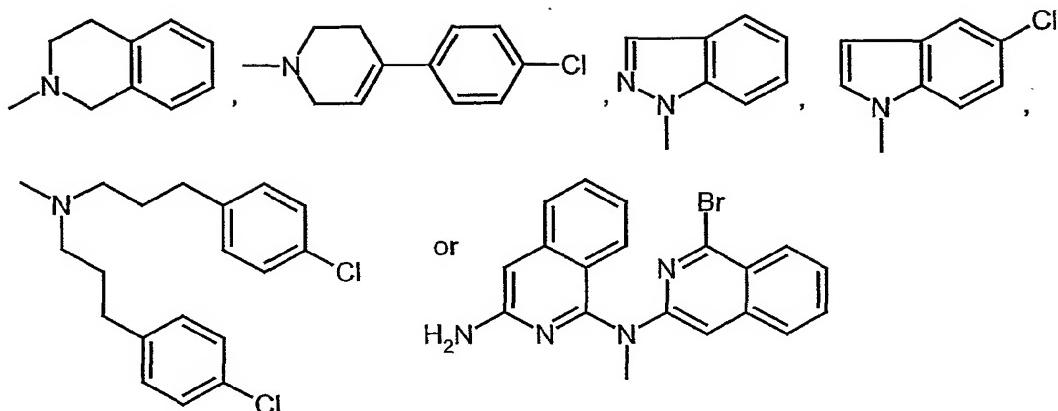
- A has the meaning of group =NR²,
- 10 W has the meaning of oxygen, sulfur, two hydrogen atoms or the group =NR⁸,
- Z has the meaning of the group =NR¹⁰ or =N-, -N(R¹⁰)-(CH₂)_q-, branched or unbranched C₁₋₆-Alkyl or is the group

15



or A, Z and R¹ together form the group

20

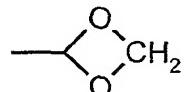


| | | |
|----|---|---|
| | m, n and o | has the meaning of 0 – 3, |
| 5 | q | has the meaning of 1 – 6, |
| | R _a , R _b , R _c , R _d , R _e , R _f | independently from each other have the meaning of hydrogen, C ₁₋₄ alkyl or the group =NR ¹⁰ , and/or R _a and/or R _b together with R _c and/or R _d or R _c together with R _e and/or R _f form a bound, or up to two of the groups R _a -R _f form a bridge with each up to 3 C-atoms with R ¹ or R ² , |
| 10 | X | has the meaning of group =NR ⁹ or =N-, |
| | Y | has the meaning of group -(CH ₂) _p , |
| | p | has the meaning of integer 1-4, |
| 15 | R ¹ | has the meaning of unsubstituted or optionally substituted with one or more of halogene, C ₁₋₆ -alkyl, or C ₁₋₆ -alkyl or C ₁₋₆ -alkoxy, which is optionally substituted by one or more of halogen, or is unsubstituted or substituted aryl or heteroaryl, |
| 20 | R ² | has the meaning of hydrogen or C ₁₋₆ -alkyl, or form a bridge with up to 3 ring atoms with R _a -R _f together with Z or R ₁ , |
| | R ³ | has the meaning of monocyclic or bicyclic aryl or heteroaryl which is unsubstituted or optionally |
| 25 | | |

5

 R^4, R^5, R^6 and R^7

substituted with one or more of für halogen, C₁₋₆-alkyl, C₁₋₆-alkoxy or hydroxy, independently from each other have the meaning of hydrogen, halogene or C₁₋₆-alkoxy, C₁₋₆-alkyl or C₁₋₆-carboxyalkyl, which are unsubstituted or optionally substituted with one or more of halogene, or R^5 and R^6 together form the group



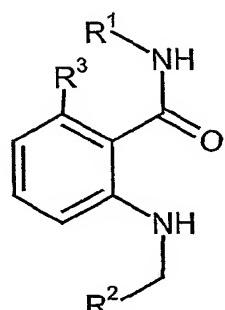
10

 R^8, R^9 and R^{10}

independently from each other have the meaning of hydrogen or C₁₋₆-alkyl, as well as their isomers and salts,

and/ or a compound of general formula V

15



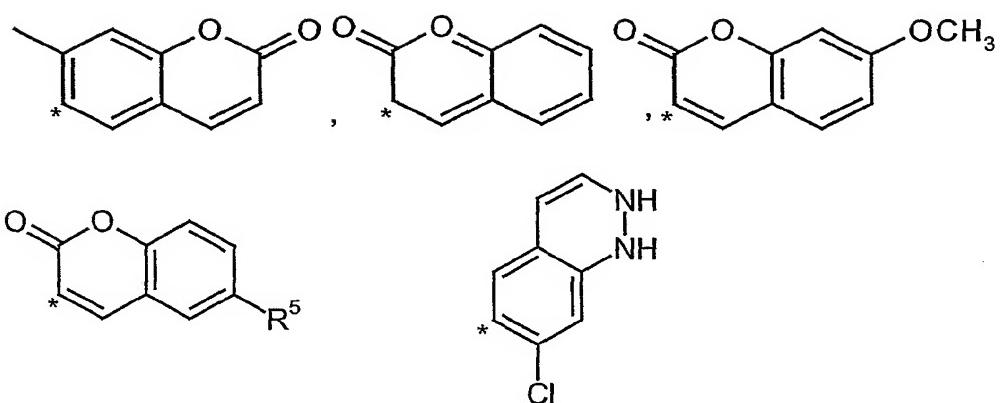
V,

20

in which

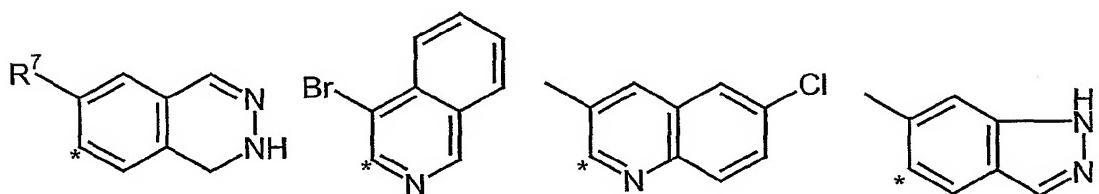
R^1 has the meaning of group

46

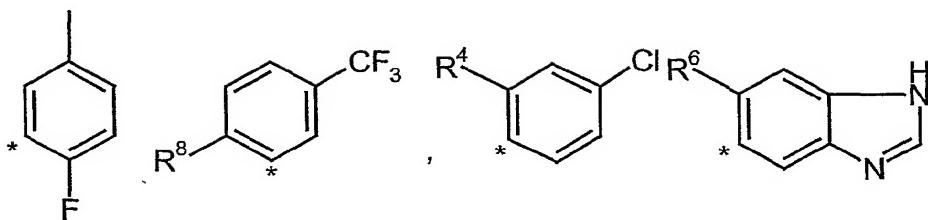


in which R⁵ is chloro, bromo or the group -OCH₃,

5



in which R⁷ is -CH₃ or chloro,



in which R⁸ is -CH₃, fluoro,
chloro or -CF₃

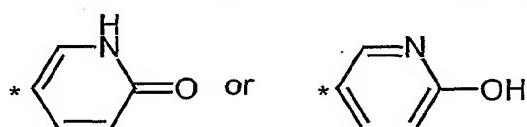
5

in which R⁴ is fluoro,
chloro, bromo, -CF₃,
-N=C, -CH₃, -OCF₃ or
-CH₂OH

in which R⁶ is
-CH₃ or chloro

R²

has the meaning of pyridyl or the group



10

and

R³

has the meaning of hydrogen or fluoro, as well as their
isomers and salts.

16. Pharmaceutical compositions according to claim 15 which comprise as

15 compound I and/ or II (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate

17. Pharmaceutical compositions according to claims 1-16 which comprise as

20 compound I (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate, sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate, and as compound II (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium hydrogen succinate, Tie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate, with the proviso that compound I is not identically to compound II.

25

18. Pharmaceutical compositions according to claims 1-17 which comprise as

compound I (4-Chlorophenyl)[4-(4-pyridylmethyl)-phthalazin-1-yl]ammonium

hydrogen succinate and as compound II sTie2, mAB 4301-42-35, scFv-tTF and/ or L19 scFv-tTF conjugate.

19. Pharmaceutical compositions according to claims 1-17 which comprise as

5 compound I mAB 4301-42-35 and as compound II sTie2, and/ or scFv-tTF conjugate.

20. Pharmaceutical compositions according to claims 1-17 which comprise as

compound I scFv-tTF conjugate and as compound II sTie2 and/ or mAB 4301-
10 42-35.

21. Pharmaceutical compositions according to claims 1-17 which comprise as

compound I L19 scFv-tTF conjugate and as compound II sTie2.

15 22. Use of pharmaceutical compositions according to claims 1-21, for the

production of a medicament for the treatment of tumors, cancers, psoriasis, arthritis, such as rheumatoide arthritis, hemangioma, angiofibroma, eye diseases, such as diabetic retinopathy, neovascular glaucoma, kidney diseases, such as glomerulonephritis, diabetic nephropathie, malignant

20 nephrosclerosis, thrombic microangiopathic syndrome, transplantation rejections and glomerulopathy, fibrotic diseases, such as cirrhotic liver, mesangial cell proliferative diseases, arteriosclerosis, damage of nerve tissues, suppression of the ascites formation in patients and suppression of VEGF oedemas.

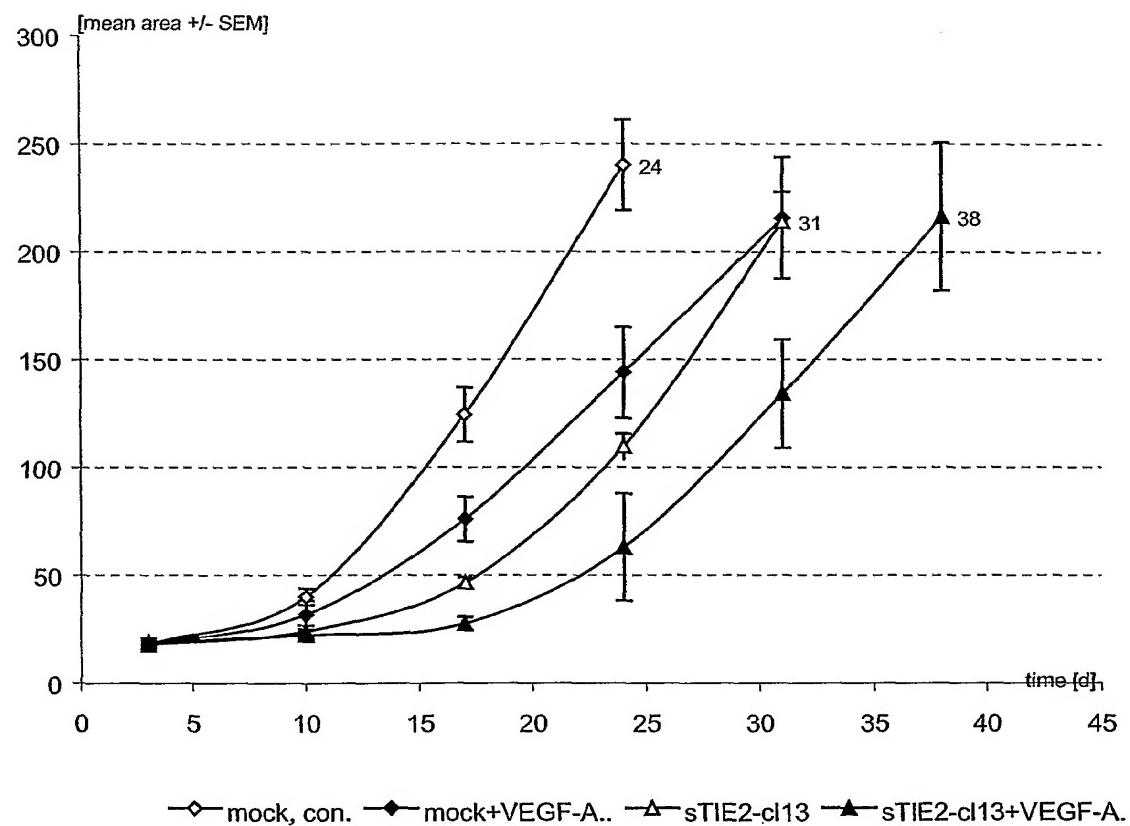


Fig. 1

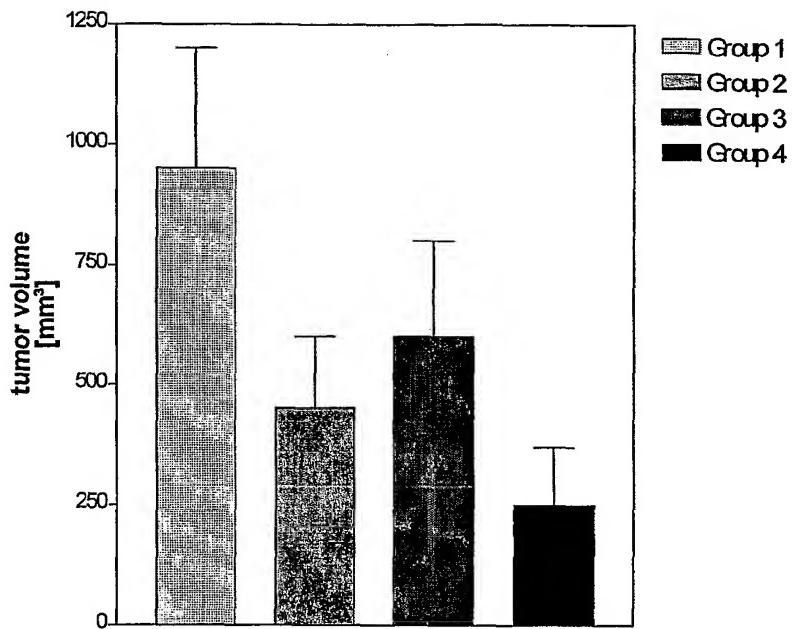


Fig. 2

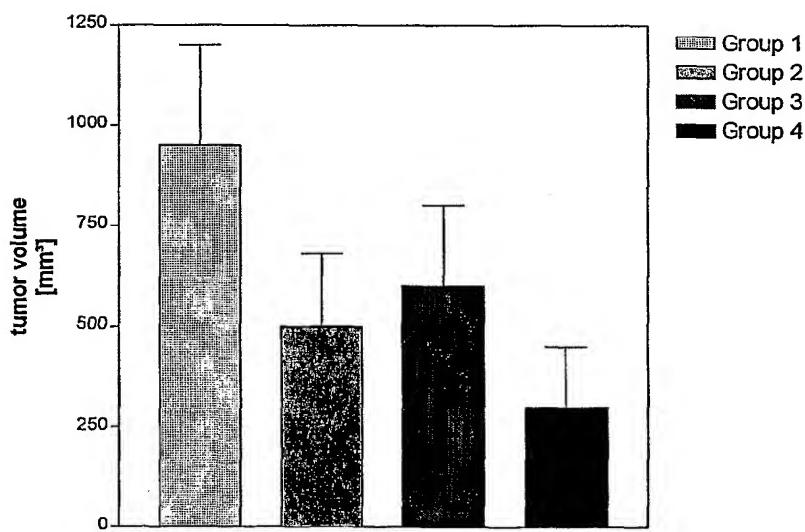


Fig. 3

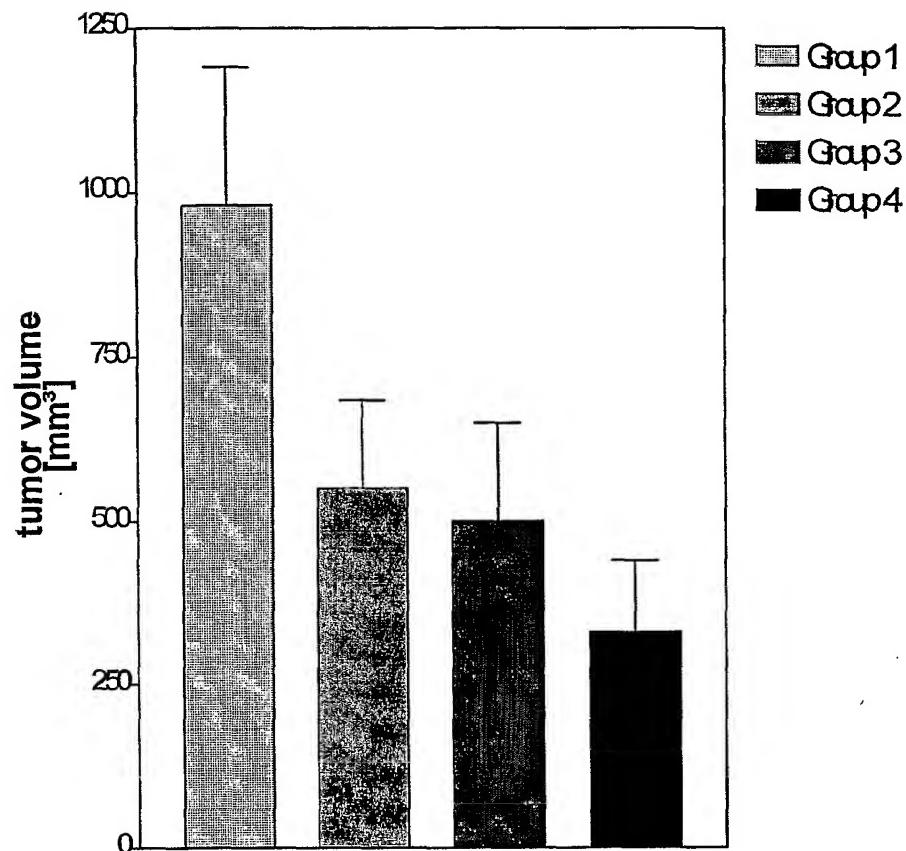


Fig. 4

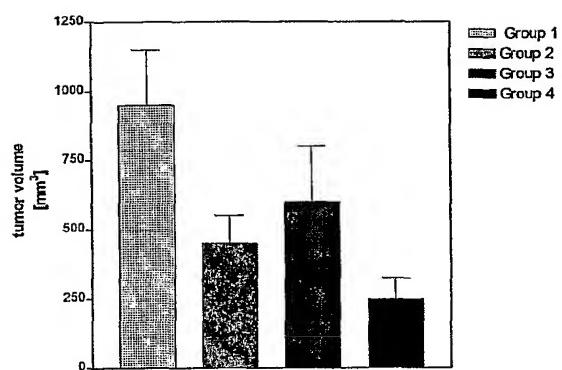


Fig. 5

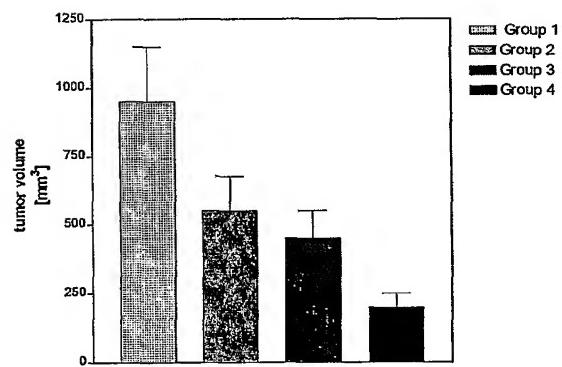


Fig. 6

Sequence Identifier

5

<110> Schering Aktiengesellschaft

10 <120> Combinations and compositions which interfere with VEGF/ VEGF and
angiopoietin/ Tie receptor function and their use II

<130> 51867AEPM1XX00-P

15 <140>
<141>

<160> 59

20 <210> 1
<211> 1835
<212> DNA
<213> Human

25 <400> 1

ttttacagtt ttccctttct tcagagttaa ttttgaattt tcatttttgataaaccaagc 60
agctctttaa gaagaatgca cagaagagtc attctggcac ttttggatag tacataagat 120
tttcttttttttttttaat agtcacattc agtcgccttgc ctcaaaccagg 180
actccccacat tgggtgagca agatgagccc ataggattcc aggttaata cgtaaccgt 240
tatacaaaca gccaaaaaaac cataatggtg ccacaggat ggagcaggga agggcatctc 300
taacgtgtcc tctagtctat cttcgctaaa cagaacccac gttacacatg ataactagag 360
agcacactgt gttgaaacga ggatgctgac cccaaatggc acttggcagc atgcagtt 420
aagcaaaaga gacatccctt aataactgta taaaatccag gcagttccat taaagggtt 480
aagaaaacca acaacaacaa aaagcgagg actgtctgtt gtcactgtca aaaaggcact 540
tggagttaat gggaccagga ttggaggact cttagctgat acagattca gtacgatttc 600
attaaaaggc ttggatgtta agagaggaca ctcagcggtt cctgaaggga gacgctgaga 660
tggaccgctg agaagcgaa cagatgaaca caaaggaatc aaatctttac aaccaaattt 720
catttaagcg acaacaaaaa aaggcaaaacc cccaaacgc acctaaccac agcaaaatct 780
aagcaaaatc agacaacgaa gcagcgatgc atagcttcc tttgagagaa cgcataccct 840
gagacgctac gtgcacact aagtctcaa cgacagcttcc acagtaggat tattgtata 900
aaaatgactc aagcgatgc aaaagttca tctttccca gaatccggg gagaactgag 960
gtgatcgatc gacatcacgtg cggtttctt atgtccctgg tggcggatc 1020
gccgagtctt cggaggaca tctggacacc accttcagcc accttccttgc agggcgaca 1080
tccgcacaaag tcatccctta ttcccgatcaa taacttaat tccttctaa catttacacg 1140
gcaaacacgga atgcagtaaa cgtccacgtc cgtcccacgg ctgggctgcc gttccgtttc 1200
ctccacgaaac gggtaacgcgc ttccatgaga aaggatattt ggcaatttttattccacag 1260
tcaggtgggt ctgcgatagc tcatttaatg ttaaacfcca tcaggggctt ctcctccgt 1320
ttctgccagg ggctttctt gtcttctctt tggcgagctc gtgggctcgtt cttctctgg 1380
gggggctggc tgctggctcc gagggggcat cccgacttcc tctggcgttgc tccttctgca 1440
ggctgggcaag ctggccacca cttctccgac tcgacccctc caacaagcat cgcaggcac 1500
tgttcctcggtt ggtacagacc gtggtcccac attcgatacc actctgttcc acgtcatcca 1560
gttacacgag ctgcgtgttag gccgtgtgtt ctggggctcg aggcttttc tgctgggtgt 1620
cttggacggg cgggttagttc tgctgcagag acaaagcatc tccccttccc ttccgggctg 1680
55 attttggttt attcatatatct acggccagatg ccaaactggc atcattactt ccgttccttc 1740
cagcttttg gagaatcaat gtatgaatgt ctaacctgac cgttggaccc gccatccaag 1800
gagacgaacc acgcccgggg gtgcggaaagc ggct

60 <210> 2
<211> 581
<212> DNA
<213> Human

<400> 2

| | |
|----|--|
| 5 | gttctagatt gtttattca gtaatttagct cttaaagaccc ctggggcctg tgctacccag 60 acactaaca cagtctctat ccagttgctg gttctgggtg acgtgatctc cccatcatga 120 tcaacttact tcctgtggcc cattagggaa gtggtgacct cgggagctat ttgcctgttg 180 agtgcacaca cctggaaaca tactgcttc atttttcat ccacatcagt gagaaatgag 240 tggcccgtta gcaagatata actatgcaat catgcaacaa agctgcctaa taacatttca 300 tttattacag gactaaaagt tcattattgt ttgtaaaagga tgaattcata acctctgcag 360 agttatagtt catacacagt tgatttccat ttataaaggc agaaaagtct tggtttctct 420 aatgtcaag ctttgactga aaactcccgt ttttccagtc actggagtggt gtgcgtatga 480 aagaaaaatct ttagcaatta gatggggagag aaggaaaata gtacttgaaa tgttaggcct 540 cacctccccca tgacatcctc catgagcctc ctgatgttagt g |
| 10 | |

15 <210> 3
<211> 516
<212> DNA
<213> Human

20 <400> 3

| | | |
|----|--|---|
| 25 | tagagatgtt ggttgatgac ccccgggatc tggagcagat gaatgaagag tctctgaaag tcagcccaga catgtgcata tacatcacag aggacatgct catgtcgccg aacctgaatg gacactctgg gttgattgtg aaagaaaattg ggtcttccac ctcgagctt tcagaaaacag ttgttaagct tcgtggccag agtactgatt ctcttccaca gactatatgt cgaaaaaccaa agacctccac tgatcgacac agcttgagcc tcgatgacat cagactttac cagaaaagact tcctgcgcacat tgcaaggcttg tgcaggaca ctgcctcagag ttacaccctt ggatgtggcc atgaactgga tgagggaaaggc ctctattgca acagggtctt ggccccagcag tgcatacaaca tccaagatgc ttttccagtc aaaagaacca gcaaatactt ttctctggat ctcactcatg 30 atqaaqtgcc aqaqtttgtt qtqtaaaagtgc cgtctg | 60 120 180 240 300 360 420 480 |
|----|--|---|

35 <210> 4
<211> 1099
<212> DNA
<213> Human

<400> 4

| | | | | | | | |
|----|--------------|-------------|-------------|-------------|-------------|-------------|------|
| 40 | cccacaaacac | aggggccctg | aaacacgcca | gccttcctc | tgtggtcagc | ttggccccagt | 60 |
| | cctgctca | ggatcacagc | ccattgttag | tggggcatgg | tggggatcag | ggcccctggc | 120 |
| | ccacggggag | gtagaagaag | acctggtccg | tgttaagggtc | tgagaagggtg | ccctgggtcg | 180 |
| | ggggtgtcg | ttggccttgc | cgtgccctca | tccccggct | gaggcagcga | cacagcaggt | 240 |
| | gcaccaaactc | cagcaggta | agcaccagg | agatgagtcc | aaccaccaac | atgaagatga | 300 |
| 45 | tgaagatgt | cttctccgtg | gggcgagaga | caaagcagtc | cacgaggtag | gggcgagggtg | 360 |
| | ctcgctggca | cacaaacacg | ggctccatgg | tccagccgt | caggcgcac | tggccataga | 420 |
| | ggaaggcgtc | ctctagcaca | ctcttgccaga | gcacactggc | gacatagtg | ccatcatgtg | 480 |
| | ctcccgccgat | gchgaggcga | ccatcttctg | ccaccgagat | cttggccatc | tgacgctcta | 540 |
| | cgccgcggccag | cgcccgctcc | acctgtgggt | ccttggccgg | cagtgcgc | agctccccct | 600 |
| 50 | ccttctgcgc | cagccgctct | tctcgccgag | acaggttaat | gacatggccc | aggtagacca | 660 |
| | gggtgggtgt | gctgacgaaag | aggaactgca | gcaccccgta | gcggatgtgg | gagatgggaa | 720 |
| | aggcctgttc | atacagacg | tttgtgcage | ctgggtggcc | cgtgttacac | tcaaatactg | 780 |
| | actgctcg | accccacact | gactcgccgg | ccaggccccag | gatgaggatg | cgaaagatga | 840 |
| | agagcaccgt | cagccagatc | ttacccacca | cggtcgagtg | cttctggacc | tggtccagca | 900 |
| 55 | acttttccac | gaagccccag | tcacccatgg | ctccccggcc | tccgtcg | aggagacaga | 960 |
| | gcacgtca | gtgtcagcat | ggcatccitc | tcgttgc | agcaacaagc | ctgcagggag | 1020 |
| | gtctgcccacg | ccctttctac | cgcctgcctg | ccggggcgcc | caggtggagg | tggggacgat | 1080 |
| | qqccqqaqtq | acqcccqcg | | | | | |

60 <210> 5
<211> 1015
<212> DNA
<213> Human

<400> 5

ccggatgggg ccgttggggg ccggatgggg ccggatgggg ccggatgggg tttccaaaaa 59

aaaaagtgt ttttggaaat gttgagggtt aatgatggg aaccaacatt ctttggattt 120
 agtggggagc ataatacgaa acacccctt ggttcgcaca tgtacagggaa tgggaccag 180
 ttggggcaca gccatggact tccceggcct ggaatgtgtg gtgcaaagtg gggccaggc 240
 ccagacccaa gaggagaggg tggtcgcag acacccggg atgtcagcat cccccgacct 300
 5 gccttctggc ggcacccccc gggtgtgt tgagtgacg aggcatgggg tgagagcctg 360
 gtatatgctg ggaacagggt gcaggggcca agcgttctc cttcagccct gacttggcc 420
 atgcacccccc tctcccccac acacaaacaaa gcacttctcc agtatggtc caggacaggt 480
 gtcccttcag tcctctgggt atgacctaagc gtcctactt ggcctgcag cccagcctgt 540
 gttgtaacct ctgcgtcctc aagaccacac ctggaagatt cttctccct ttgaaggaga 600
 10 atcatcattt tgcgttatac acttctaaga cattttgtac ggcacggaca agttaaacag 660
 aatgtgttcc cctccctggg gtctcacacg ctcccacag aatgccacag gggccgtgca 720
 ctgggcaggc ttctctgttag aaccccgagg gcttcggccc agaccacagc gtcttgcct 780
 gaggcttagag cagggagtcc cgaacttctg cattcacaga ccacctccac aatttgtata 840
 15 accaaaggcc tcctgttctg ttatctact taaatcaaca tgctatttt ttttactca 900
 cttctgactt tagcctcgtg ctgagccgt tatccatgca gtcatgttca cgtgttagtt 960
 acgtttttct tcttacacat gaaaataaaat gcataagtgt tagaagaaaa aaaaa

<210> 6
 <211> 2313

20 <212> DNA
 <213> Human

<400> 6

25 ccagagcagg ectgggtggt agcagggacg gtgcacccga cggcgggatc gagcaaatgg 60
 gtctggccat ggagcacgga gggtcctacg ctcgggggg gggcagctt cggggctgct 120
 ggtattacct ggcgttacttc ttccttctcg tctccctcat caaatccttc atcatcctgg 180
 ggctcgtgt cttcatggtc tatggcaacg tgcacgtgag cacagagttc aacctgcagg 240
 30 ccaccgagcg ccgagccgag ggcctataca gtcagcttctt aagggtcactg gcctccctgt 300
 ccaacttgcac caaggagctc aacttcacca cccgcgcacaa ggatgcattc atgcagatgt 360
 ggctgaatgc tcgcgcgcac ctggaccgcac tcaatgcacag cttccgcac tgccagggtg 420
 accgggtcat ctacacgaac aatcagaggt acatggctgc catcatctt agtgagaagc 480
 aatgcagaga tcaattcaag gacatgaaca agagctgcga tgccttgcac ttcatgtca 540
 atcagaagggt gaagacgctg gaggtggaga tagccaagga gaagaccatt tgcactaagg 600
 35 ataaggaaag cgtgtgtgt aacaaacgcg tggcggagga acagctgggtt gaatgcgtga 660
 aaacccggga gctgoagcac caagagcgc actggccaag gagcaactgc aaaaggtgca 720
 agccctctgc ctgcccctgg acaaggacaa gtttgagatg gaccttcgtt acctgtggag 780
 ggactccat atccacgcac gcctggacaa cctgggttac aacctctacc atccctctggg 840
 40 ctcggattt gcctccatcc gcagggcctg cgaccacatg cccagcttca tgagctccaa 900
 ggtggaggag ctggccggga gcctccgggc ggatatecgaa cgcgtggccc gcgagaactc 960
 agacctccaa cgccagaagc tggaaagccca gcagggtctt cggggccatgc aggaggcgaa 1020
 acagaagggtt gagaaggagg ctcaggcccg ggaggccaag ctccaaagctg aatgtcccg 1080
 gcagaccccg cttagcgtgg aggagaaggc ggtgtcgcc aaggaacgag acaacctggc 1140
 45 caaggagctg gaagagaaga agaggggggc ggagcagctc aggtatggagc tggccatcag 1200
 aaactcagcc ctggacaccc tcatcaagac caagtcgcag ccgtatgtc cagtgtcaag 1260
 gcccattgggc cctgtccccca acccccaagcc catcgaccca gtcagcttgg aggagttcaa 1320
 gaggaagatc ctggagtccc agaggcccccc tgcaggcatc cttgttagccc catccagtgg 1380
 ctgaggaggt tccaggcctg aggaccaagg gatggcccgat ctcggcggtt tgccggaggat 1440
 50 gcagggatata gtcacacgcg cccgacacaaa cccctccctt cgcggccatggg ccacccagg 1500
 ccaccatcag acaactccct gcatgcaaaac cccttagtacc ctctcacacc cgcaccccg 1560
 ctcacacgc cctcacccctc agcacaacggc cggggagatg acgtcacgc agcaacggcg 1620
 ctgacgtcac atatcaccgt ggtgatggcg tcacgtggcc atgttagacgt cacgaagaga 1680
 tatacgatg gctgtgtca gatgcagcac gtcgcacaca gacatggggaa acttggcatg 1740
 55 acgtcacacc gagatgcagc aacgcgtca cggggcatgt cgacgtcaca cataattaatg 1800
 tcacacagac gcccgcgtatgg catcacacag acgggtatgt tgcacacac agacacagt 1860
 acaacacaca ccatgacaac gacacctata gatatggcac caacatcaca tgcacgcatt 1920
 cccttccaca cacactttt acccaattt cacttagtgc cacgttcccc cgaccctggc 1980
 acacggggcca aggtacccac aggtacccat cccctcccg acagccctgg gccccagcac 2040
 60 ctcccctctt ccagcttccctt ggcctcccg ccacttctc acccccaatgt cctggaccccg 2100
 gaggtgagaa caggaagcca ttcacctccg ctccttgagc gtgagtgttt ccagggcc 2160
 ctggggccctc tgagccgggg gtgagggtca cctgtgtcg ggaggggagc cactccttct 2220
 cccccaactc ccagccctgc ctgtggcccg ttgaaatgtt ggtggcactt aataaatatt 2280
 agtaaatcct taaaaaaaaaaa aaaaaaaaaaaa aaa

65 <210> 7
 <211> 389

<212> DNA
<213> Human

<400> 7

5 gccaaaaaga tggcttcaa agtaagaatg aaacatttga tccattcagc tttaggctat 60
gccactggat tcatgtctag aaaagataagg ataatttctg taaagaaaatg aagaccttgc 120
tattctaaaa tcagatcctt acagatccag atttcaggaa acaaatacat agggactaa 180
10 ctttccttgt tcagattagt ttttccttgc tgaccccagc tatataatat gaggaagtat 240
tgactttta aaagtgtttt agtttccat ttcttgata tgaaaagtaa tatttcggga 300
gaacccttag ctattaataa tctatgtggc tagtgcgtat atattggct gaatttggc 360
tcctttgtg gtgtccagtg ggtaaacatc

<210> 8

15 <211> 157
<212> DNA
<213> Human

<400> 8

20 tgctttaaac agctgtgtca aaaactgaca tcagagagta aattgaattt ggttttgttag 60
gaagcagggaa gcaagccac tcaaacgtga aatttggcat gaggatcca gtaactttct 120
cctcaatctg tgaactatataa gtgagtttga tattttg

25 <210> 9

<211> 561
<212> DNA
<213> Human

30 <400> 9

aatagtcaaa acataaacaa aagctaattt actggcactg ttgtcacctg agactaagtg 60
gatgttgttgcgtt gctgacatac aggctcagcc agcagagaaaa gaattctgaa ttcccttgc 120
35 tgaactgaac tattctgtt catatggttt acaaattctgt gtgttatttc ttttctacct 180
accatatttta aatttatgag tatcaaccga ggacatagtc aaaccttcga tgatgaacat 240
tcctgatttt ttgcctgatt aatctctttt gggctact tttttttttt caagattttt 300
tgatgttggaa aggaaaagtg aatatgaccc ttaaaaatttga tattttttttt gatgatagtc 360
tcaccactat aaaactgtca attattgctt aatgtttaaag atatccatca ttgtgattaa 420
40 ttaaaccttat aatgagtattt ctaatggag aattcttaat ggtatggatca tccctgtatc 480
ttttttttttt aatttctctg cacacacagg acttcttcatt ttccaataaaa tgggtgtact 540
ctggcccaat ttcttagaaaa a

<210> 10

45 <211> 1508
<212> DNA
<213> Human

<400> 10

50 cacaacacag agagactcca cggtctgcctt gggcccgcc agcctcctag gctccagcac 60
tcgcagggtcc attttctgc acgagccctt ctgtccagat ccataaggcac ggtcaagctca 120
gggtcgccga gcagtagcgg gacaagtacc agcagcagct cctctgaaca gagactgctca 180
55 ggatcatctt tctccctccgg gcctgttgcgtt gatggatcaaa tccgggtgc acccaaatct 240
gagctcaagc caggtagct taagccactg agcaaggaag atttggccct gcacgcctac 300
aggtgtgagg actgtggcaaa gtcaaatgtt aaggagtgc tttttttttt ccctcccaag gcctctgcca 360
tcagactgaa tctcgacaa gcagtcctt tgctcggccc agaacgtgtat tgactatggg 420
acttgtgtat gctgtgtgaa aggtctcttc tatactgtt ctaatgtatga tgaggacaac 480
tgcgtgtaca acccatgttc ttgcagccag tctcactgtt gtacacgtat gtcagccatg 540
60 ggtgtcatgt ccctttttt gccttgcattt tggtgttacc ttccagccaa gggttgcctt 600
aaattgtgcc aggggtgttgcgtt aacaggcctg gttgcgcgtt taaaaactca 660
aacacagtt gctgaaatgtt tccactgtc ccccttagga actttgaaaa accaacatag 720
catcatataat caggaatattt acagtaatgtt ggttttttcc ttctttttt taatcacat 780
atgcaaccaa ctaaacagttt ataatcttgg cactgtttaat agaaaagttgg gatagtctt 840
65 gctgtttgcgtt gtttttttttgcgtttt aactgtatgtt cttttttttt gttttttttt 900
ctcagctaat ggagctcaaa gtatggatata cagaacttgg tgaccatgtt attgcataag 960
ctaaagcaac acagacactc cttaggcaaaat ttttttttt tgaatagtac ttgcacaaact 1020

tgtaaatttag cagatgactt ttttccattg ttttctccag agagaatgtg ctatatttt 1080
gtatatacaa taatatttg aactgtaaa aacaagtggt gccatactac atggcacaga 1140
cacaaaatat tatactaata tggtgtacat tcggaagaat gtgaatcaat cagtatgttt 1200
tttagattgtt tttgcctta cagaaagcct ttattgtaaag actctgattt cccttggac 1260
ttcatgtata ttgtacagtt acagtaaaaat tcaacctta ttttctaatt ttttcaacat 1320
attgttttagt gtaaaaagaata ttatttgaa gttttattat ttataaaaaa agaatattt 1380
tttaagagg catcttacaa attttggccc ttatgtagg atgtgatagt tgctgcaaat 1440
gaggggttac agatgcatat gtccaaata aatagaaaa tatattaacg ttgaaattt 1500
aaaaaaaaaa

10 <210> 11
<211> 389
<212> DNA
<213> Human

15 <400> 11

20 gggcaggta tcagggcaca catttccgt ccattgagac agtagcattc ccggcaccca 60
tcgtgccagc ttccttcatt ttatgtatga tgaccatcca cggtgagaca agtgcggac 120
aggatgggtg gcccagctga agcacaggcc gctctgcact tgcaagataag acagccgtga 180
ctgtcctgtt ggaaacccaa ggggcagatc ttactgcatt agagctctgg acatttttta 240
cagcgacaga tgtcacagcc gtgcttattc ttcagcaatc caagtggaca atacttgtca 300
cagattatgg gtctgcactt cttggccctt gggccggact cacagatctc acagttttgg 360
acctcqccq cqaccacgct qqgtaccqa

25 <210> 12
<211> 981
<212> DNA
<213> Human

30 <215> Human
<400> 12

| | |
|----|---|
| | ttttttttt ttggattgca aaaatttatt aaaattggag acactgttt aatcttcttg 60 |
| 35 | tgcctatgaga ctccatcagg cagtctacaa agaccactgg gaggctgagg atcacttgag 120 |
| | cccagaagtt tgaggctgta gtaagcttca aaggccactg cactctagct tgggtgaggc 180 |
| | aagacccttt caagcagtaa gctgcattgtc tgcttgggt ggtcattaaa aacccttagtt 240 |
| | taggataaca acatattaat cagggcaaaa tacaaatgtg tgatgcttgt tagtagagta 300 |
| 40 | acctcagaat caaaatggaa cgggtttaca gtgatatcat tatatttcat ttggcagaat 360 |
| | cattacatca ttggttacac taaaaatcat cacatgtacc aaaagctgac tcacctagtt 420 |
| | taggataaca ggtctgcctg tttgaagatg aaaaataata cccatTTaaa atttgcccta 480 |
| | ctcaatttcc ttctoagtca catttttaact tttaaacagc taatcactcc catctacaga 540 |
| | ttaagggtgtt tatgccacca aaaccttttgc ccaccttaaa aatttccttc aaagttaaa 600 |
| | ctaatgcctg cattttctca atcatgaatt ctgagtcctt tgcttcttta aaacttgctc 660 |
| 45 | cacacagtgt agtcaagccg actctccata cccaaagcaag tcatccatgg ataaaaacgt 720 |
| | taccaggagc agaaacattt agetggtcca ggcaaggttgg actccacat ttcaacttcc 780 |
| | agctttctgt ctaatgcctg tttggccaaatg gcttggatgtt ggcttgctt ttaggactt 840 |
| | agtagctatt ctcatccctt cttggggaca caactgttca taaggtgtca tccagagcca 900 |
| | cactgcattt gcaccccgca ccatacctca caggagtcga ctccccacgag ccggctgttat 960 |
| | ataagagtcc ttttggatqac q |

50 <210> 13
<211> 401
<212> DNA
<213> Human

55 <400> 13

ataactacag cttcagcaga caactaaaga gactgcatta aggtgatttc tctggctata 60
 aagagagccc ggccgcagag catgtgactg ctggacctx tggataggc aacactgcc 120
 tctctcccccc agagcgaccc cccgggcagg tcggggccca aggaatgacc cagcaactgc 180
 tccctaccca gcacactctc tttactgc当地 cctgcaattt tgctgtgaag atgactgggt 240
 gtggtcatca cgattcagag aaatcaagat ctatgaccat ttttaggcaaa gagagaaaact 300
 tggagaattt ctgaggacta ctgaaccttg ttttgctttt taaaaaaaata ctaaatcctc 360
 acttcagcat atttagttgt cattaaaatt aagctqatat t

65

<211> 1002

<212> DNA

<213> Human

5 <400> 14

gacaatataa aaagtggaaa caagcataaa ttgcagacat aaaataatct tctggtagaa 60
acagttgtgg agaacagggtt gagtagagca acaacaacaa aagcttatgc agtcacctc 120
tttggaaaatg ttaaatacaa gtcctattct ctttgtccag ctgggttttag cttagaggtag 180
10 ccaattactt ctcttaaggt ccatggcatt cgccaggatt ctataaaagc caagtttaact 240
gaagtaaata tctggggccc atcgaccccc cactaagtagc tttgtcacca tgggttatct 300
taaaagtcat ttttactgt ttgactcaga atttggact tcagagtcaa acttcattgc 360
ttactccaaa cccagttta cttcccaccc ttttaagtag gtttagctt gaggatgtt 420
15 tggctataac cgaatgtaa atccaccttc aaacaacaaa gtttgacaag actgaaatgt 480
tactgaaaac aatggtgcca tatgctccaa agacatttcc ccaagataac tgccaaagag 540
tttttgagga ggacaatgtat catttattat gttaggagcct tgatatatct gcaaaataga 600
attaatacag ctc当地atgga gtagtaacca agctttctg cccaggaatg aacaaacatc 660
actacgaaca tgagagtaca agaggaaact ttcataatgc atttttcat tcatacatc 720
attcaataaa cattagccaa gctaattgtcc caagccactg tgccaggtat taacaatata 780
20 acaacaataa aagacacagt ctttccttc aaggtttca gtcttagtagg gaagatgatt 840
attcattaaa atttttgggtg catcagaatc atgaggagct tgtcaaaaaat gtaaattcct 900
gcctatgttc tcagatattc tggtaggtc aggagtggga accccaaaatc aattctttta 960
acaaacacta aaggtgattc taacacaggc ggtgtgagga cc

25 <210> 15

<211> 280

<212> DNA

<213> Human

30 <400> 15

cgagggtggc caccctgttc tggctctgaga tttttaatg aggattacat tatcctattt 60
ataatattcc tattttaatc tattgtattt ttacaattaa atgtatcaaa taattcttaa 120
35 aaacattatt agaaacaaac tgcctaatac cttataagac taaaaaaaaatc accaagatga 180
actgttatta tgactctcaa tatttaaaca tttaaaaaaaaa tggtagtgtt tggtagcac 240
caatcttaac tatttcaccc gccccggcgg ccgctcgagg

<210> 16

<211> 2041

<212> DNA

<213> Human

<400> 16

45 ccccccgcag aactcccccc tggaaatagga tttttaaaac ctttgcacat tagaaatcct 60
atagaggta gcattttta ggtaaaaata tggtgcccc tacagggatc atgcaacttc 120
cttaaaacca attcagcaca tatgtataaa gaacccttt taaaaacatt tggtagtgc 180
atacagacac agtggatgtc aagacactaa acaaaaactg aaaagtacta taccttgata 240
aattttgtta ttgccttctt tagagacttt ataatctcta gttgattttc aaggacttgc 300
50 atttaataat ggggttaatta cacaagacgt aaaggatttt taaaaacaa gtatTTTTT 360
ttacctctag catcaattct ttatataaga atgcttaataa aattacatt tttgttcagt 420
aaaactgaag atagaccatt taaaatgttc taccaaaattt aacgcagctt aatttagggac 480
caggtacata ttttcttcg aacatttttgc tcaagcatt tctaaccata aaagcaaatg 540
gaatttttaag aggttagattt ttttccatg atgcattttgc ttaataaaatg tggtagaaaa 600
55 ataaaaacaa gcactgagtg tggctcttcg aagtataagg gtctaatgaa aaataaaaga 660
tagatatttg ttatagtcgt acatttttaac agtcatagta ttagacgtt cgtgaccagt 720
gcattttgga ctctctcagg atcaaaatac gagtctgccaa actgtattaa atccctcc 780
acccccctcca ccagggttc cacagcttcc tgggggtcg ttgtcatcaa atccattggg 840
ccgaaatgaa catgaagcag atgcagtttgc gaggcccgg gctcgagcat tcaactctt 900
60 ttcctgtaaa tatagttat tggctttgt tatagcatcc ataagttttc tctgttagagg 960
tgggtctcca ttatccaga gtccactgtt tgggttattt ccacttaaac cattagact 1020
atgctgtttt ttatataaaa gcacataagc tggctctttt gggaaacctgc tcgtatTTTT 1080
ctggactgac tggaaatgtcaatgtca tctactgtca taaaataaaa acccattttt 1140
ttgacatttc cttatTTTCC aaatcctgtt caaaaactgc actgggacta tctctcccta 1200
65 gtaaatgact ctggaggat gctaatgc gaggctcaga ctgggttgc atctgtatgt 1260
aagagtctgtt acttgcatt aagaatagta atgcccactt tcagaggata 1320

taccagagt aaccacaacg gaacttaata gatagggcac caattttgtg caggaagctt 1380
 catca~~gt~~ccc tgaaggctt aatttttag caaggtctc actaagatca gtgaagtcaa 1440
 catctacaga ccaactttct gacaatgaag agaaaagaat aatttctcta actggcaact 1500
 cccaaaccag tggccagtga tacattgtct aaaatttcc ttctcacatg atacttctga 1560
 5 tcata~~tg~~aaa atctcaggag agtaagaata aggtattcag gttcctccgt gatttgcata 1620
 gtttctcag cattttgcag agaggcacag ttttacaat aatattggtt atcaccagta 1680
 agaatctctg gagccaaaaa aataatttag taagt~~c~~agtt actgaagg~~tg~~ tggtttcacc 1740
 tccccgtt~~c~~ tgaggtacat ctttattaac aagaatctt~~g~~ ttagattcgt tagggacaga 1800
 10 agt~~ttt~~tca gaacagtaaa actcattagg aggactgcct atgg~~ttt~~ttt cattcacaag 1860
 tgagt~~c~~acag atgaaggcag ctgtt~~ttt~~gg attataaact actggctt~~c~~ ctgaaggacc 1920
 ggt~~t~~acagac gctgcatt~~a~~ gaccaccatc ttgtatactg ggtgatgat~~g~~ ctggatctt~~g~~ 1980
 gacagacat~~g~~ tttccaaag aagaggaagc acaaaacgca agcgaaagat ctgtaaaggc 2040
 t

15 <210> 17
 <211> 235
 <212> DNA
 <213> Human

20 <400> 17

cgccccgggc aggtgtcagg ggttccaaac cagcctgggg aaacacagcg tagacccctc 60
 acctctacaa ataaaaaaatt aaaaaattag ccaggtgtgg cagcgaacaa ctgt~~at~~gtctc 120
 25 agatactcag gagactgagc tggaaaggat cacttgagcc caagaagttc aagg~~t~~acag 180
 tggccacaga tcatgtcatt acactccagc ttgggtgaca aaatgagact gtcta
 <210> 18
 <211> 2732
 <212> DNA
 30 <213> Human

<400> 18

gtgtggagtt tcagctgcta ttgactataa gagctatgga acagaaaaag cttgctggct 60
 35 tcatgtt~~g~~at aactactt~~a~~ tatggagctt cattggac~~c~~ gttac~~c~~ttca ttattctgt~~c~~ 120
 aatattttac ttcttgg~~t~~ga tcacattgt~~g~~ caaaatgg~~t~~ aagcatt~~c~~aa acactttgaa 180
 accagattct agcagg~~t~~gg aaaa~~c~~attaa gtcttgg~~t~~g cttggcgc~~t~~ tcgc~~c~~t~~c~~t~~c~~ 240
 gtgt~~c~~t~~c~~t~~c~~ ggc~~c~~t~~c~~ac~~c~~ ggtt~~c~~ttttt~~t~~ attaatgagg agactattgt 300
 40 gatggcatat ct~~c~~t~~c~~t~~c~~acta tatttaatgc tttccagg~~g~~ gttt~~c~~ttt~~c~~tc~~c~~at~~c~~ttt~~c~~ca 360
 ctgtgctctc caaaagaaag tacgaaaaga~~a~~ atatggcaag tgctt~~c~~agac actcatactg 420
 ctgtggag~~g~~ c~~t~~cccaactg ag~~g~~t~~c~~ccca cagtt~~c~~ag~~t~~ aaggcat~~c~~aa ccaccagaac 480
 cagt~~c~~t~~c~~gc tattc~~c~~t~~c~~g~~c~~acacagag tcgtataaga agaatgtg~~g~~ atgatactgt 540
 gaaaaa~~c~~aa tcagaatctt ctttat~~c~~tc aggt~~g~~acat~~c~~ aatagcactt caacacttaa 600
 tcaagg~~t~~gg~~c~~ ataaatctt atatatttt acaggact~~g~~ catcacat~~g~~ tctgagagcc 660
 45 cat~~c~~tt~~c~~aa~~g~~ atttat~~t~~ca ttt~~c~~agg~~g~~ac attcact~~g~~aa caat~~g~~ccagg gatacaag~~t~~g 720
 ccat~~g~~gata~~c~~ tctacc~~g~~cta aat~~g~~taatt ttaacaacag ctactcg~~c~~t~~c~~acaagg~~g~~t~~g~~ 780
 actataat~~g~~ a~~c~~gc~~g~~t~~g~~caa gtt~~t~~ggact~~g~~ gtggact~~g~~taag tctgaat~~g~~at actg~~c~~ttt~~g~~ 840
 50 agaaaaat~~g~~at catticagaa tt~~g~~at~~g~~caca acaacttac~~g~~ gggcag~~g~~c~~g~~ aagactcaca 900
 acctcgag~~c~~ c~~a~~c~~g~~ct~~c~~acca g~~t~~caa~~a~~c~~t~~g~~c~~ tgatt~~g~~gagg tagcag~~c~~agt gaagat~~g~~atg~~g~~ 960
 ctattt~~g~~tg~~c~~ agat~~g~~t~~c~~ta tttt~~a~~tgc acag~~g~~aca~~a~~ cccagg~~g~~ct~~g~~ gag~~c~~ccat~~c~~ 1020
 acaaagaact cgagg~~g~~acca cttatt~~c~~tc~~c~~ a~~g~~cg~~g~~act~~c~~a cttcc~~c~~tt~~c~~g tacc~~a~~cccc 1080
 agaagaaaa~~g~~at ga~~g~~t~~c~~cg~~g~~ g~~g~~aa~~g~~t~~g~~aca gctat~~g~~t~~c~~tc~~c~~aa~~g~~t~~g~~aca gcagagg~~g~~ct~~g~~ 1140
 aagat~~c~~ac~~c~~t ac~~g~~t~~c~~cccc aac~~g~~ag~~g~~act ct~~c~~t~~c~~t~~c~~ata aagcat~~g~~ccc aat~~c~~ttagag 1200
 55 actctcc~~c~~ta tccggagagc agcc~~c~~t~~g~~aca tgg~~a~~gaaga~~g~~ cctct~~c~~ccc tccaggagga 1260
 gtgagaat~~g~~ta ggacatttac tataaaag~~g~~ca tgccaaat~~c~~t tggag~~c~~tt~~g~~gc catca~~g~~ctt~~c~~ 1320
 agat~~g~~t~~g~~cta ccagat~~c~~gc~~c~~ aggg~~g~~caata g~~t~~gat~~g~~g~~t~~ta tataat~~c~~ccc attaaca~~a~~ag 1380
 aagg~~g~~tg~~t~~at tccagaagga gat~~g~~tagag aaggacaat~~g~~ g~~c~~ag~~g~~t~~g~~g~~t~~tt acaag~~t~~ctt~~c~~ 1440
 aatcata~~c~~ag ctaagg~~a~~att ccaagg~~g~~cca cat~~g~~cg~~g~~at~~c~~ ttaataaata aagacaccat 1500
 60 tggc~~c~~t~~g~~ac~~g~~ cag~~c~~cc~~c~~tc~~c~~ aaact~~c~~t~~g~~ct~~c~~ tgaagagat~~g~~ act~~c~~tt~~g~~acc~~c~~ t~~g~~tg~~t~~tc~~c~~ 1560
 tgg~~t~~gt~~aaaa~~ aagat~~g~~act~~g~~ aac~~c~~tt~~g~~ca~~g~~ tt~~c~~t~~g~~gaat ttttataaaa catacaaaaa 1620
 ct~~t~~t~~g~~t~~t~~at~~at~~ acacag~~g~~act~~g~~ tactaa~~g~~at~~g~~ aatttattt~~g~~t~~c~~ taca~~a~~agaaa agagat~~g~~cc~~a~~ 1680
 g~~c~~cagg~~g~~t~~t~~ ttaagg~~a~~t~~c~~ g~~c~~t~~g~~t~~g~~tt agaaaatt~~g~~tg~~a~~aca~~g~~cc~~a~~aaaacaaaac 1740
 tt~~t~~t~~c~~ag~~g~~cca tt~~t~~t~~c~~act~~g~~ca g~~c~~ag~~g~~t~~c~~tg~~g~~ aactaaat~~g~~tt~~a~~at~~g~~tg~~c~~tc~~g~~acc~~at~~t~~g~~ 1800
 tt~~t~~t~~g~~agg~~g~~cc t~~g~~catt~~g~~tgat tataaca~~g~~ag~~g~~ ac~~g~~t~~g~~agg~~g~~tt taaaat~~c~~ct~~g~~ tgg~~g~~acaat~~g~~ 1860
 65 tt~~t~~act~~g~~t~~acc~~ tt~~t~~act~~t~~ttcc tgacaagact tgg~~a~~aaag~~g~~ca ggagagat~~g~~at tct~~g~~cat~~g~~ 1920
 tt~~t~~tg~~c~~agt~~tc~~ act~~g~~caa~~a~~at~~c~~ tt~~t~~tacat~~g~~ta agg~~c~~aaag~~g~~at tgg~~a~~aaagg~~g~~at cttaacc~~act~~ 1980

agcaatcaag ccacaggcct tatttcataat gtttcctcaa ctgtacaatg aactattctc 2040
 atgaaaaatg gctaagaaa ttatatttg ttcttattgct agggtaaat aaatacattt 2100
 gtgtccaact gaaatataat tgtcattaaa ataattttaa agagtgaaga aaatattgtg 2160
 aaaagcttt gggtgcacat gttatgaaat gtttttctt acactttgtc atggtaagtt 2220
 5 ctactcatt tcacttctt tccactgtat acagtgttct gctttgacaa agttagtctt 2280
 tattacttac atttaaattt cttattgcca aaagaacgtg ttttatggg agaaacaaac 2340
 tcttgaagc cagttatgtc atgccttgca caaaagtgtat gaaatctaga aaagattgtg 2400
 tgtcacccct gtttattctt gaacagaggg caaagaggc actgggact tctcacaac 2460
 tttctagtga acaaaggtg cttattctt ttaaaaaaaaaaaa cataaataattt 2520
 10 actcttccat attccttctg cttatatttta gtaattaattt tattttatga taaagttcta 2580
 atgaaatgt aattgtttca gaaaattct gctttttt catcccttg tgtaaacctg 2640
 ttaataatgt gccccatcact aatatccagt gtaaagtttta acacggttt acagtaaata 2700
 aatgtgaatt ttttcaagtt aaaaaaaaaaa aa

 15 <210> 19
 <211> 276
 <212> DNA
 <213> Human

 20 <400> 19

 ctcccataat gattttaaaa taaattggat aaacatatga tataaagtgg gtactttaga 60
 aaccgcctt gcatatttt tatgtacaaa tctttgtata caattccgat gttccttata 120
 25 tattccctat atagcaaacc aaaaccaggc cctcccaact gcatgcctca agtccctgtg 180
 gagcactctg gcaactggat ggcctactt gcttctgac aaaatagctg gaaaggagga 240
 gggaccaatt aaatacctcg gccgcgacca cgctgg

 <210> 20
 <211> 2361
 30 <212> DNA
 <213> Human

 <400> 20

 35 attgtaccag cttgtatgaa cgtggccct gttcgctt tgagggccat aagtcattt 60
 cccactgggt tagaggctac ttatcattt tctccgtt ccggaaagggt tctcccaagt 120
 cagagtttac cagcaggat tcacagact ccgacaagca gattctaaac atctatgacc 180
 tgtgcaacaa gttcatacgcc tatagcacgg tcttggagga ttagtggat gtgcggctg 240
 40 agtggggctc cttgtacgtt ctgcacggg atggggggg ccacgcactg caggagaagg 300
 acacacagac caaactggat atgttgttta agaagaacactt atttgatgat gcgattaaacc 360
 ttgccaagag ccagcatctg gagactgtat ggctggccca gattttcatg cagttggag 420
 accatctcta cagcaagggc aaccacgtt gggctgttca gcaatatact cgaaccattt 480
 gaaagttgga gccatccatc gtgatccgca agtttctgga tgcccagcgc attcacaacc 540
 tgactgccta cctgcagacc ctgcaccggc aatccctggc caatgcgcac cataccaccc 600
 45 tgctcctcaa ctgtatacc aagctcaagg acagtcgaa gctggaggag ttcatcaaga 660
 aaaagagtga gagtgaagtc cactttgtat tggagacagc catcaaggatc ctccggcagg 720
 ctggctacta ctccccatgcc ctgtatctgg cggagaacca tgacatcat gatgtgtacc 780
 tgaagatcca gctagaagac attaagaattt atcaggaagc ctttcgatc atcggcaagc 840
 tgccttttga gcaggcagag agcaacatga agcgtacgg caagatctc atgcaccaca 900
 50 taccagagac gacaactcg ttgctgaagg gactttgtac tgattatcg cccagcctcg 960
 aaggccgcac cgatagggg gcccagggt gcaggccaa ctctgaggag ttcatcccc 1020
 tctttccaa taacccgcga gagctgaaag ctttcctaga gcacatgatgaaatgcgc 1080
 cagactcacc ccaggggatc tacgacacac tccttgcgt ggcactgcac aactggggcc 1140
 acgagaagaa tccacaggc aaagagaagc ttacgcgcaga ggcctatttc ctgtgaaga 1200
 55 gtggtcgtt ctgcacgtc ttgtacaagg cccttgcctt gtgcgcacatg cacgacttcc 1260
 aggatgggtt ctttacattt tatgagcagg ggaagctgtt ccagcagatc atgcactacc 1320
 acatgcagca cgagcgtac cggcagggtca tcagcgtgtg tgagcgcacat ggggaggcagg 1380
 acccctcctt gtggggcag gcccctcagatc acttcgctcg caaggaggag gactgcaagg 1440
 60 agtatgtggc agctgtcctc aagcatatcg agaacaagaa cctcatgcca cctttcttag 1500
 tgggtgcagac cttggccac aactccacag ccacactctc cgtcatcagg gactacctgg 1560
 tccaaaaact acagaaacag agccagcaga ttgcacagga tgagctgcgg gtgcggcgg 1620
 accgagagga gaccacccgt atccgcacgg agatccaaga gctcaaggcc agtcttaaga 1680
 ttttccaaa gaccaagtgc agcatctgtt acagtgcctt ggagttgccc tcagtccact 1740
 tctgtgtgg ccacttcctc caccaacact gtttgcgtt gttactcgaa agtgcgtctg 1800
 65 actgccccac ctgcctccctt gaaaaccggg aggtcatgga tatgatccgg gcccaggaaac 1860
 agaaacgaga tctccatgat caattccacg atcagctcaa gtgcgtccaaat gacagctttt 1920

ctgtgattgc tgactacttt ggcagagggtg tttcaacaa attgactctg ctgaccgacc 1980
 ctcccacagc cagactgacc tccagcctgg aggctggct gcaacgcgac ctactcatgc 2040
 actccaggag gggcaactaa gcagcctgga ggaagatgtg ggcaacagtg gaggaccaag 2100
 agaacagaca caatgggacc tggcgggcg ttacacagaa ggctggctg catgccagg 2160
 5 gctccactct catctaatgt cacagccctc acaagactaa agcggaaacctt tttctttcc 2220
 ctggccttcc ttaattttaa gtcaagcttg gcaatccctt cctcttaac taggcagggtg 2280
 ttagaatcat ttccagatta atggggggga agggaaacctt caggcaaacc tcctgaagtt 2340
 ttggaaaaaaa aagctggttt c

10 <210> 21
 <211> 179
 <212> DNA
 <213> Human

15 <400> 21

aggtgttaga tgctctttagaa aaagaaaactg catctaagct gtcagaaaatg gattctttta 60
 acaatcaact aaaggaactg agagaaaacct acaacacaca gcagttagcc cttgaacagc 120
 20 tttataagat caacgtgaca agttgaagga aattgaaagg aaaaaatttag aactaatgc
 <210> 22
 <211> 905
 <212> DNA
 <213> Human

25 <400> 22

tttttttttt ttctttaacc gtgtggtctt tatttcagtg ccagtgttac agatacaaca 60
 30 caaatgttcc agttagaaaaagg aattcaaaacg gaatgccaag gtccaaagcca ggctcaagaa 120
 ataaaaaaggg aggtttggag taatagataa gatgactcca atactcaactc ttcttaaggg 180
 caaaggtaact ttgatacag agtctgatct ttgaaacttgg tgaactccctc ttccacccat 240
 taccatagtt caaacaggca agttatgggc tttaggagcac tttaaaattt gtggtggaa 300
 tagggtcatt aataactatg aatatatctt tttagaagggtg accatttgc actttaaagg 360
 gaatcaattt tgaaaatcat ggagactatt catgactaca gctaaagaat ggcgagaaag 420
 35 gggagctgaa agaggcttgg aagtttctat tacaatataa gcaccatatac ttcatgcca 480
 aatctcaaca aaagctcttt ttaactccat ctgtccagtg tttacaataa aactcgcaag 540
 gtctgaccag ttcttggtaa caaacatatac tttgtgtgtc tttgtgtata cagcaatgca 600
 cagaaaaaggc taccaggagc ctaatgcctc ttccaaacat tgggggaacc agtagaaaaa 660
 40 ggcagggtctc cctaattgtcc atttacat ttccattccg aatgccagat gttaaaagtg 720
 cctgaagatg gtaacccttgc tagtgaggaa taaatacccc accttgccca gtccacagag 780
 aaacaacagt agaaaagaagg ggcaacttgc tgctgcagag acaaagttag tttttttcg 840
 ccatggattt cagttctctc ttccagacca gctgttattt tcctcagggg cccaggaaat 900
 gttga

45 <210> 23
 <211> 2134
 <212> DNA
 <213> Human

50 <400> 23

ggtctcttct ttccaaaaat ttttccaaa agtgttcttt tatttcagtg aacatataatt 60
 gtataaaatc tctattttat atgcacttcc acaaaaagcgataataattttaa aagttttttt 120
 55 cattagaaaat aaatgtataa aaataaaatattt gttattatag gcattttatta ctaactatag 180
 tccttcttgg aaggaacacc caaaccataa cttataaagt acatgtattt tatagtaaca 240
 tattttacta tatacatatg gaaaaatca tatttcacca gaagagctga acagacattc 300
 accaggatac gactgttggc ccagctgctg gagatggacc tgctaccctc cagcagccctc 360
 cccaccacaa gacaagtgtat ctaatgtcc ccaaaacctgt gggaccctgt tctacacacc 420
 tcatttttgt tccggcgttt catcccttgc ttgtgtattt actgattttc atgagacaca 480
 60 agttacttct ttacatccat attcccaaag cagggttaca ttgttagggaa gaaaggaaat 540
 tggaggtact aagctcattt gttctcttgc agcttttacc agcatctaat gcttcactgc 600
 ttttttttca ttgttagactt taatgcactt gaataaaatc atggagttgt tttttctca 660
 aatgaattt cacaataaa gactgagatg gtccaaaaaaa gggaaagagga agccatttgc 720
 65 gttatttcac gttgtgagc ctttctctca ttgttgacaa tctgaagttt taattctcg 780
 tagaaataat gtataaaacat tctctgaaac catagcagcc ataaacagtgc ctggtcaaag 840
 atccttatttgc tactcctttc tccccccattt gtttagtgggg taaagtaaaa caggctttag 900

taaaatctca cttttctcct acttttcatt tcccaaccccc catgatacta agtatttgat 960
 aagtaccagg aaacagggggt tgtaatagtt ctaactttt ttgacaattg ctttgggtt 1020
 tctaaacttg taatagatgt aacaaaagaa ataataataa taatgccccc ggcttatta 1080
 5 tgctatatca ctgctcagag gttataatc ctcactaact atcctatcaa atttgcact 1140
 ggcagttac tctgatgatt caactcctt tctatctacc cccataatcc caccttactg 1200
 atacaccccta ctgggtactg gcaagatacg ctggatccct ccagccttct tgcttccct 1260
 gcaccagccc ttccctcactt tgccttgcct tcaaagctaa caccacttaa accacttaac 1320
 tgcattctgc cattgtgcaa aagtctatga aatgttttagg tttctttaaa ggatcacagc 1380
 10 tctcatgaga taacaccccct ccatcatggg acagacactt caagcttctt ttttgtaa 1440
 cttcccccaca ggtcttagaa catgatgacc actcccccaag ctgcccactgg gggcagggat 1500
 ggtctgcaca aggtctgggt ctggctggct tcacttcctt tgccactctg gaagcaggct 1560
 gtccattaat gtctcggcat tctaccatc ttctctgcca acccaattca catgacttag 1620
 aacattcggcc ccactcttca atgaccatcg ctgaaaaagt ggggatagca ttgaaagatt 1680
 15 cttcttctt ctttacgaag tagtgtatt taatttttagg tgaaggggca ttgcccacag 1740
 taagaacctg gatggtcaag ggctttga gaggctaaa gctgcgaattt ctttccaatg 1800
 ccgcagagga gccgcgtgtac ctcaagacaa caccttgcataatgtct tgctctaagg 1860
 tggacaaaagt gtagtcacca ttaagaatat atgtccatc agcagcttg atggcaagaa 1920
 agctgccatt gttctggat cccctctgt tccgtgtt cacttcgatg ttggtggctc 1980
 20 cagttggaat tgtgatgata tcatgatatac caggtttgc actagtaact gatcctgata 2040
 ttttttaca agtagatcca ttccccccgaa acaccaca ttatcaa 2100
 agtctatgat gcgatcacaa ccagtttta caca

<210> 24
 <211> 1626
 25 <212> DNA
 <213> Human

<400> 24

30 ggacaatttc tagaatctat agtagtatca ggatataattt tgctttaaaa tatattttgg 60
 ttatttgaa tacagacatt ggctccaaat tttcatctt gcacaatagt atgactttc 120
 actagaacctt ctcaacattt gggactttt caaatatgag catcatatgt gttaaaggctg 180
 tattatattaa tgctatgaga tacattgttt tctccctatg ccaaacaggt gaacaaacgt 240
 35 agttgtttt tactgatact aaatgttggc tacctgtat tttatagttt gcacatgtca 300
 gaaaaaggca agacaaaatgg cctcttgcac tgaataacttc ggaaacttta ttgggtcttc 360
 attttctgac agacaggatt tgactcaata ttttagagc ttgcgtagaa tggattacat 420
 ggtagtgtat cactggtaga aatggttttt agttatttgc tcagaattca tctcaggatg 480
 aatcttttat gtcttttat tgaaggcata tctgattttt ctttataaag atggtttttag 540
 40 aaagctttgt ctaaaaattt ggccttaggaa tggtaacttc attttcagtt gccaaggggat 600
 agaaaaataaa tatgtgtt gttatgtt ttttacata ttatttaggtt ctatctatga 660
 atgtattttttaa atattttca tatttcgttca caagcattttt taatttgcac caagtggagt 720
 ccatttagcc cagttggaaa gcttggac tcagttacc ttttgcattttaa agtgcggcag 780
 ccatctctt gatctgttca taaactgtttaa ttttagtacc agctaaatcc ctaacttgaa 840
 45 tctggaatgc attagttatg ctttgcacca ttcccaaaat ttcaggggca tcgtgggtt 900
 ggtctagtga ttgaaaacac aagaacagag agatccagct gaaaaagagt gatcctcaat 960
 atcctaacta actggcctc aactcaagca gagtttctt actctggcac ttttgcatttgc 1020
 aaacttagta gaggggattt tttttttttaa atacaatttt aatacaatgt ttttgcatttgc 1080
 taaaattctt aaagagcaaa actgcattttt ttttgcatttgc ttttgcatttgc 1140
 50 aactaagata tttatctatg aagatataaa tggtgccagag agactttcat ctgtggattt 1200
 ctttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1260
 atggatttttctt ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1320
 atgcccaggatg ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1380
 ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1440
 55 ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1500
 attataacctg tcacgcttctt ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1560
 tacttggaaaa ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc ttttgcatttgc 1620
 aaaaaa

60 <210> 25
 <211> 1420
 <212> DNA
 <213> Human

<400> 25

gttcagcatt gtttctgctt ctgaaatctg tatagtacac tggtttgc ttttgcatttgc 60

| | | | | | | |
|-------------|------------|-------------|-------------|-------------|-------------|------|
| ttcattgtaaa | tccttgctac | ttcttccct | cctcaatgaa | agacacgaga | gacaagagcg | 120 |
| acacaagctt | aagaaaaacg | agcaaggaag | agtatcttca | ttattctcat | tttctctgag | 180 |
| ttggaaacaa | aaacatgaag | gactccaact | agaagacaga | tatttacatt | taaatagatt | 240 |
| agtggaaaaa | ctttaagagt | ttccacatata | tagtttcat | tttttgagtc | aagagactgc | 300 |
| tccttgtact | gggagacact | agtagtatata | gtttgtatag | ttacttaaa | attatcttt | 360 |
| tatTTtataa | ggcccataaa | tactggtaa | actctgttaa | aagtgggcct | tctatcttgg | 420 |
| atggttcac | tgcctatcgc | catgtgtata | tatttagaaa | ggcatcccta | tctacttact | 480 |
| ttaatgtta | aaattataca | taaaatgttt | tatttagaaa | acctacatga | taagtggtg | 540 |
| tcagccttgc | catgtatcag | tttcaacttga | aatttgagac | caattaaatt | tcaactgtt | 600 |
| agggtggaga | aagaggtact | ggaaaacatg | cagatgagga | tatctttat | gtgcaacagt | 660 |
| atccttgca | tgggaggaga | gttacttctt | aaaggcaggc | agcttaagtg | gacaatgtt | 720 |
| tgtatatagt | tgagaatttt | acgacacttt | taaaaattgt | gtaattgtt | aatgtccagt | 780 |
| tttgccttgt | tttgcctgaa | gttttagtat | ttgttttctt | ggtgacacc | tgaaaaccaa | 840 |
| accagtacct | ggggagggtt | gatgtgttt | tcaggcttg | agtgtatgag | tggttttgc | 900 |
| tgtattttcc | tccagagatt | ttgaacttta | ataattgcgt | gtgtgtttt | tttttttttaa | 960 |
| gtggctttgt | ttttttttct | caagtaaaaat | tgtgaacata | tttcctttat | aggggcaggg | 1020 |
| catgagttag | ggagactgaa | gagtattgtt | gactgtacat | gtgccttctt | aatgtgtttc | 1080 |
| tcgacacatt | ttttttcagt | aacttgaaaa | ttcaaaaaggg | acatctgggtt | aggttactgt | 1140 |
| acatcaatct | atgcataaat | ggcagcttgt | tttttttgagc | cactgtctaa | attttgggtt | 1200 |
| tatagaaaatt | tttatactg | attgggtcat | agatggtca | ttttgtacac | agactgaaaca | 1260 |
| atacagact | ttgccaaaaa | tgagtgtac | attgtttaa | cattgtgtgt | taacacctgt | 1320 |
| tctttgttaat | tggggtgtgg | tgcattttgc | actacacgtt | tttacagttt | tcaatctgtc | 1380 |
| agtaaataaa | gtgtccttta | acttcaaaaa | aaaaaaaaaa | | | |

25 <210> 26
<211> 689
<212> DNA
<213> Human

30 <400> 26

```

aaacaacaaa aaaaaaagtt agtactgtat atgtaaatac tagctttca atgtgctata 60
caaacaatta tagcacatcc ttccctttac tctgtctcac ctcctttagg tgagtacttc 120
cttaaataag tgctaaacat acatatacgg aacctgaaag ctgggttag ccttgcctta 180
ggtaatcagc ctatgttaca ctgtttccag ggagtagttt aattactata aaccattagc 240
caacttgtctc tgaccattt atcacaccag gacagggctc ctcaacctgg gcgtactgt 300
cathttgggc cagggtattc ttcccttgaa gggctgtcct gtacctgccc gggcgccgc 360
tcgaaggctg gtcgcggccg aggtactgaa aggaccaagg agctctggc gcccttcagga 420
attccaaatg accgaaggaa caaaggttca gggctctgg tggtgcctc cactattcag 480
gagggtgtcg gaggttaacgc agcttcaattt cgttcaggctc ttccctttagat tttaagggtt 540
tgtcaagatg ctgcattaaa tcaggcagg tctacaaaggc atccccaaagca tcaaacatgt 600
ctgtgtatgaa gtaatcaatg aaacaccggaa acctccgacc acctcctgaa tagtgggaga 660
cacaccggaa gcctgaagtt tgtcctcg

```

45 <210> 27
<211> 471
<212> DNA
<213> Human

50 <400> 27

| | | | | | | |
|-------------|------------|------------|-------------|-------------|------------|-----|
| tcccagcggc | atgaagtttg | agattggcca | ggccctgtac | ctgggcttca | ttccttcgt | 60 |
| ccctctcgct | cattgggtgc | accctgctt | gcctgtccctg | ccaggacgag | gcaccctaca | 120 |
| agcccctaacc | caggccccgc | ccagggccac | cacgaccact | gcaaaacacgg | cacctgccta | 180 |
| ccagccacca | gctqcctaca | aqacaatcq | qqccccctca | gtgacacctgg | ccaccacagc | 240 |
| gggtacaggc | tgaacgacta | cgtgtgagtc | cccacagcct | gcttctcccc | tgggctgctg | 300 |
| tgggctggtt | cccgccggga | ctgtcaatgg | aggcaggggt | tccagcacaa | agtttacttc | 360 |
| tgggcaattt | ttgtatccaa | gaaaataatg | tgaatgcgag | gaaatgtctt | tagagcacag | 420 |
| ggacagaggg | gaaaataaga | ggaggagaaa | gctctctata | ccaaagactg | a | |

60
<210> 28
<211> 929
<212> DNA

```

ggtaactca gtgcattggg ccaatggtc gacacaggct ctgccagcca caaccatcc 60
gctgtttctg acggtttggc tgctgggg cttccccctc actgtcattg gaggcatctt 120
tggaaagaac aacgcagcc ccttgatgc accctgtcgc accaagaaca tcgccccgg 180
gattccaccc cagccctggt acaagtctac tgtcatccac atgactgtt gagggttcct 240
gccttcagt gccatctctg tggagctgta ctacatctt gccacagtat ggggtcggg 300
gcagtagtact ttgtacggca tcctcttctt tgcttcgccc atcctgtga gtgtggggc 360
ttgcatctcc attgcactca cctacttcca gttgtctggg gaggattacc gctgggtgg 420
gcatctgtg ctgagtgtt gctccacccg ccttcatac ttccctactt cagttttcta 480
ttatgccccg cgctccaaca tgctggggc agtacagaca gtagaggctt tcggctact 540
cttactact ggttatgtct tcttcctcat gctgggcacc attcctttt ttcttcctt 600
aaagtttcatc cggatatct atgttaacct caagatggac tgatgtctgt atggcagaac 660
tattgctgtt ctccctt cttcatgccc tggtaactc tcctaccagc ttcttcctctg 720
attgactgaa ttgtgtatg gcatgttgc ctcccttt tcccttggg catttccttc 780
ccagagaggg cctggaaatt ataaaatctt atcacataag gattatatat ttaaactttt 840
taagttgcct ttagtttgg tcctgtttt tcttttaca attacaaaaaaa taaaattttat 900
taaaaaaaaaa aaaaaaaaaa aaaaaaaaaa

```

20 <210> 29
<211> 1775
<212> DNA
<213> Human

29

| | |
|----|---|
| 25 | gaacgtgatg ggaactttgg gaggatgtct gagaaaaatgt ccgaaggat tttggccaac 60 accagaaaac gccaatgtcc taggaattcc ctccccaaat gtttcccaaa aaattactca 120 ttgacaattc aaattgcact tggctggcgg cagcccgccg ggccttcagt ccgtgtgggg 180 cgcccgcgtg gccttcttc cgttaggactc cccaaactcg ttcactctgc gtttatccac 240 30 aggataaaagc caccgttgtt acaggatgac cagaaacac acgtcgccc ggaagcaggc 300 cagccgtga gacgtggca tggtgatgat gaaggcaaag acgtcatcaa tgaagggtt 360 gaaaggcttg taggttaagg cttccaggg catatgtgcc actgacttca acttgtatg 420 cacaagagc tgggcagca tgaagaggaa accaaaggca tagaccctgt tgacgaagct 480 gttattttttt tttttttttt ttttttttttt ttttttttttt ttttttttttt ttttttttttt 540 cccgacacag agagggtaca gcaggtatga caagtacttc atggccttag tatctactc 600 35 ctcggttttc ctctcagatt cgctgttaagt gccaaactga aattcggca tcaggccct 660 ccaaaaaata gtcatttca atgccttctt cacttccac agctcaatgg cggctccaac 720 acccgcccgg accagcacca gcaggctcg ctgctgtcc agcaggaaca gaaagatgac 780 40 cacggtgctg aagcagcgcc agagcactgc cttgggtggac atgcccgtca tgctttctt 840 cttcttccag aaactgtatgt catttttaaa ggccaggaaa tcaaagagaa gatggAACgc 900 tgcacaaag aagggtcagcg ccagagaatg taagtgttca tctacaaaaaa ttccatttcac 960 45 ctcatcagca tctttttctg aaaacccggaa ctgctgcagg gaggatcacgg cgtcttcac 1020 gtggatccag aagcgcagcc gccccagtg gaccttgcg taggacacgg tgaggggcag 1080 ctcggtgggt gaggcggtta tgaccatcg gtccttcacg cgggtgtcga gctggcgt 1140 gaacaggatg ggcaggtaat gcacggttt ccccaagctgg atcatttca tgtaccgatg 1200 cacatcggtca ggcaggggagg acccgtaaaa gacaaggatg tccgccccatca cgttcagcgc 1260 cagccgcggg cgccagtggg acactggctc atccaggcga ctgcgtggct tcttcggc 1320 50 ctcgatctgc tgtgtatcag actccccggt gaggcggtt gatcttcgt gcttggggac 1380 catgttaggtg gtccaggaggac tgaccaggtg caccctgttc ccgtgtgcc acggcaggac 1440 cccagcgtga tggaggaaga tgtaggcata cagcgtccca ttgtttctcg ttttttttgg 1500 tacagaaaaca ttaactgtcc ttccaaattt ggactccaca tcaaagtctt ccacattcaa 1560 gaccaggctcg atgttgttct cagccaccccg gtgggacctc gtctgtgtgtt acacgctcag 1620 55 ctgcgttgtt gggccggccgg ccaggtaggg ctggatgcag ttggcgctcgc cggagcacgg 1680 gcgggtgttag acgatggcgtt acatgaccca gcagggtgtc accacgtaga ccacgaacac 1740 acccaccacc aagctgtga aggagctggc gcccc |
|----|---|

<210> 30
<211> 1546
<212> DNA
60 <213> Human

<400> 30

aaaataagta ggaatggca gtgggtattc acattcacta cacctttcc atttgctaat 60
aaggccctgc caggctggga gggaaattgtc cctgcctgct tctggagaaaa gaagatattg 120

| | | | | | | | |
|----|-------------|--------------|--------------|-------------|--------------|--------------|------|
| | acaccatcta | cgggaccat | ggaactgctt | caagtgacca | ttcttttct | tctgcccagt | 180 |
| | atttcagcga | gtAACAGCAC | agggtttta | gaggcagcta | ataattcaact | tgttgttact | 240 |
| | acaacaaaac | catctataac | aacacccaaac | acagaatcat | tacagaaaaaa | tgttgtcaca | 300 |
| | ccaacaactg | gaacaactcc | taaaggaaca | atcaccaatg | aattactaa | aatgtctctg | 360 |
| 5 | atgtcaacag | ctacttttt | aacaagtaaa | gatgaaggat | tgaaagccac | aaccactgat | 420 |
| | gtcaggaaga | atgactccat | catttcaaac | gtaacagtaa | caagtgttac | acttccaaat | 480 |
| | gctgttcaa | cattacaaag | ttccaaaccc | aagactgaaa | ctcagagttc | aattaaaaca | 540 |
| | acagaaatac | caggtgtgt | tctacaacca | gatgcacat | cttctaaaac | tggtacatta | 600 |
| 10 | acctcaatac | cagttacaac | tccagaaaaac | acctcacat | ctcaagtaat | aggcactgag | 660 |
| | ggtgtaaaaa | atgcaagcac | ttcagcaacc | agccggctt | atcccagtat | tatttgcgc | 720 |
| | gtgggttattt | ctttgcattgt | aataacactt | tcagtttttg | ttctgggg | tttgtaccga | 780 |
| | atgtgttgg | aggcagatcc | gggcacacca | gaaaatggaa | atgatcaacc | tcagttctgtat | 840 |
| | aaagagagcg | tgaagtttct | taccgttaag | acaatttctc | atgagtctgg | tgagcactt | 900 |
| 15 | gcacaaggaa | aaaccaagaa | ctgacagctt | gaggaattct | ctcccacacct | aggcataata | 960 |
| | taacgttaat | cttcagcttc | tatgcaccaa | gcgtggaaaa | ggagaaagtc | ctgcagaatac | 1020 |
| | aatcccgact | tccatcacctg | ctgctggact | gtaccagacg | tctgtccag | taaagtgtat | 1080 |
| | tccagctgac | atgcaataat | ttgatggaaat | caaaaagaac | cccggggctc | tcctgttctc | 1140 |
| 20 | tcacatttaa | aaattccatt | actccattta | caggagcgtt | ccttagaaaa | ggaatttttag | 1200 |
| | gaggagaatt | tgtgagcagt | gaatctgaca | gcccaggagg | tgggctcgct | gataggcatg | 1260 |
| | actttccta | atgtttaaag | tttccgggc | caagaatttt | tatccatgaa | gactttccta | 1320 |
| | cttttctcgg | tgttcttata | ttacctactg | tttagtattta | ttgtttacca | ctatgttaat | 1380 |
| | gcagggaaaaa | gttgcacgtg | tattattaaa | tattaggtag | aaatcatacc | atgtactttt | 1440 |
| | gtacatataa | gtattttatt | cctgcttcg | tgttactttt | aataaataac | tactgtactc | 1500 |
| 25 | aatactctaa | aaatactata | acatgactgt | gaaaatggca | aaaaaaaa | | |
| | <210> | 31 | | | | | |
| | <211> | 750 | | | | | |
| | <212> | DNA | | | | | |
| | <213> | Human | | | | | |
| 30 | | | | | | | |
| | <400> | 31 | | | | | |
| | cacttggcga | cccccatttt | ctaaaaaaaaat | ggaaatctgg | agggcaaaaaa | agggtgtgctg | 60 |
| | aagggaagtg | cctctgtatgg | ccaaaaaaacc | ttcttccaaa | ctagtgtagg | aatggaatgg | 120 |
| 35 | atagcaaatg | gatccctttt | ggcctccctt | ggagcatgccc | ttccctatct | tatccttggc | 180 |
| | cccactaaag | cagaacgtta | cggatatttc | tgtttttgccc | attggatgcc | tatctggcca | 240 |
| | aacagccctt | ccctaattgg | aaaatgcagt | cctgtttaaa | acctttgatt | tacgactact | 300 |
| | tgtacatgt | tgctcattac | aattttgaca | ttttttacat | agtgaagacc | ccaaacatat | 360 |
| | cagtgaaaca | tgacaagatc | ataaagaaca | gtatcatatt | attattttatgt | cgcttttaca | 420 |
| 40 | gtggcaagcc | aattttgaaa | tatctcattt | aaaactcaga | cccaatttcac | tgagtttatac | 480 |
| | tttaataatgc | ttcctcagca | cacttattcc | catgcattaa | atatgataaa | ataatctatc | 540 |
| | actcccccac | ggtcttgc | aaaggaagtc | tgaatacaga | gcccacaaaa | ctaaaattgt | 600 |
| | tttttctatgt | acaatgtata | gcatcatcaa | cacagacacg | atttggactc | cctgacaggt | 660 |
| 45 | gatttggaaa | acgggtttta | aagagaagag | aacattttaa | cataaatgtc | attaagaatc | 720 |
| | ccaaaggcct | tatttgcac | caccgtcccc | | | | |
| | <210> | 32 | | | | | |
| | <211> | 1620 | | | | | |
| | <212> | DNA | | | | | |
| | <213> | Human | | | | | |
| 50 | | | | | | | |
| | <400> | 32 | | | | | |
| | gcaattcccc | cctcccaacta | aacgactccc | agtaattatg | tttacaaccc | attggatgca | 60 |
| | gtgcagccat | tcataagaac | cttggtgc | ccagaaaaatc | tgtccttttt | ggtaccaaac | 120 |
| 55 | ctgagggttt | ttggaaagata | atgtagaaaaa | ccactaccta | ttgaaggcct | gttttggcta | 180 |
| | atctgtgcaa | actctgtatga | tacctgc | atgtggattc | tttccacac | tgttttcatt | 240 |
| | ttaatgtata | aagactttaga | aaactagaat | aatgttttta | caaataatta | aaagtatgt | 300 |
| | atgttctgg | ttttttccct | tttttagaa | ccccgcctcc | attttaaaaaa | ttaaaaaaaa | 360 |
| 60 | aaaaaaaaact | ttaacat | aaaaataaa | atttacaaa | atttcactta | ttccaggaca | 420 |
| | cgctggcatt | tggactcaat | gaaaaggggca | cctaaagaaa | ataaggctga | ctgaatgttt | 480 |
| | tccataattt | tcacacacaata | acgtccctt | tctatccagc | ttgccttcca | tttatctcta | 540 |
| | gggttagctt | ttcaggcaac | atccttggtc | attgcccaga | aagtacatgt | gctatcagt | 600 |
| | attggaatgg | cacagggaaac | cgaatcacat | gggtgc | cccttgggtt | tcaagtatct | 660 |
| 65 | tggagttgt | cacaaaaatt | aggtcatgcc | ttcagttct | ttttttttaa | acccatccctt | 720 |
| | tgacaatcag | gtgctaatga | ttgtatacta | ttaaaaaccag | cacataagta | ttgttaatgt | 780 |

gtgttcctcc taggttggaa gaaatgtct tccttctatc tgggtcctgt taaagcgggt 840
 gtcagttgtg tcttttacc tcgattttgt aattaataga attggggggga gagggaaatga 900
 tgatgtcaat taagtttcag gtttgcattg atcatcatc tcgatgatat tctactttg 960
 5 tcgcaaatct gcccttatcg taagaacaag tttcagaatt tccctccac tatacgactc 1020
 cagtattatg tttacaatcc attggatgag tgcagcatta taagacctt gtcggccagaa 1080
 aaatctgtcc ttttttgtac caaaccttagt gtcttttggaa agataatgt aaaaaccact 1140
 acctatttggaa ggcctgtttt ggtaatctg tgcaactt gatgataacct gcttatgtgg 1200
 attctttcc acactgctt catttttaag tataaagact tagaaaaacta gaataatgct 1260
 tttacaataa attaaaaggta ttttgatgttc tgggtttttt ctttctttt agaaccctgt 1320
 10 atttaacaaa gccttctttt taagtcttgg ttgaaattta agtctcagat cttctggata 1380
 ccaaatacaaa aacccaacgc gtaaaacagg gcagttttt tttttcttaat tttaaaaggc 1440
 tttatgtata ctctataat atagatgtt aaacaacact tccccttgg tagcacatca 1500
 acatacagca ttgtacatttta caatgaaaat gtgtactt agggtattat atatataat 1560
 acatatataac ctttgcatttgg tttatactgt aaataaaaaa gttgttttag tcaaaaaaaaa 1620
 15
 <210> 33
 <211> 2968
 <212> DNA
 <213> Human
 20 <400> 33

 gaaaaaagtag aagggaaacac agttcatata gaagtaaaaag aaaaccctga agaggaggag 60
 25 gaggaggaag aagaggaaga agaagatgaa gaaagtgaag aggaggagga agaggaggga 120
 gaaagtgaag gcagtgaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 180
 gggaaagacat tagataaaaaa gccaagttaa gaaatggatctt cttttttttt gatggaaaagg 240
 gatgtatgtc ggactaaaga agaaagggtt tatgacaaag cttttttttt gatggaaaagg 300
 cggcgacttg aacatagtaa aaatgttaac accggaaaagg tttttttttt gatggaaaagg 360
 gtacttgggc atgtggacac agggaaagaca aaaattctttt gatggaaaagg tttttttttt gatggaaaagg 420
 30 gtacaagatg gtgaagcagg tttttttttt caacaaattt gggccaccaa tttttttttt gatggaaaagg 480
 gaagctatta atgaacagac taagatgatt tttttttttt atagagagaa tttttttttt gatggaaaagg 540
 ccaggaatgc taatttttttta tactccttggg catgaatctt tttttttttt gatggaaaagg 600
 ggaagctctc ttttttttttta tgccattttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 660
 cagacaatgtt agtctatcaa cttttttttt tttttttttt gatggaaaagg tttttttttt gatggaaaagg 720
 35 aataagatgtt atagggttata tgatttttttta aagagtctt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 780
 ttaaaaaggc agaaaaaggaa tacaaaagat gttttttttt gatggaaaagg tttttttttt gatggaaaagg 840
 gttagatttg cacagcaggg tttttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 900
 actttttttt gttttttttt gttttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 960
 40 taccttcttgg tagagtttac tttttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1020
 ctgagagcac aggtgtatgtt gttttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1080
 atcttgcatttca atggggctttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1140
 cccatttttttta cttttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1200
 aagaaccagt atgaaaaggca taaagaagta gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1260
 45 aaagacacttgg agaaaaacattt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1320
 atccctgttc ttaaaagatgtt atttttttttt gatggaaaagg tttttttttt gatggaaaagg 1380
 tttagaaagaaa aagggttctt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1440
 gaattttctt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1500
 aaagatgtt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1560
 50 gccttcgtt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1620
 atttttttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1680
 tacaagaaaac agaaaacaaga agaaatgtt gttttttttt gatggaaaagg tttttttttt gatggaaaagg 1740
 atccctccctt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1800
 gcaggttcagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1860
 55 gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1920
 gaagtttggat gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 1980
 tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2040
 gactgggttca gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2100
 gtatgttggaa tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2160
 60 gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2220
 gacttaatgtt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2280
 aacttacactt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2340
 actcacctt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2400
 cccaaatttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2460
 gtactgtttt gttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2520
 65 acgtaagaaa tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2580
 tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg tttttttttt gatggaaaagg 2640

5
agctgctttg tggaaacca tgggtaaaaa gcacagctgg ctgccttta ctgccttgt 2700
agtcacgagt ccattgtaat catcacaatt ctaaaccaaa ctaccaataa agaaaacaga 2760
catccaccag taagcaagct ctgttaggtc tccatggta gtggtagctt ctctcccaca 2820
agttgtcctc ctaggacaag gaattatctt aacaaactaa actatccatc acactacctt 2880
gttatgccag cacctggta acagtaggat attttataca ttaatctgtat ctgtttaatc 2940
tgatcggtt agtagagatt ttatacat

<210> 34
<211> 6011
10 <212> DNA
<213> Human

<400> 34

15

20 acggggcgcc ggacgacccg cacatcttat cctccacgcc ccactcgcac tcggagcggg 60
accgccccgg actccccctc gggccggcca ctgcaggagt gaggagagag gccggccggcc 120
cggttgagc cgagcgcagc accccccggc ccccgccca gaagtttgggt tgaaccgggc 180
tgccgggaga aactttttc tttttccccc ctctcccggt agagtctctg gaggaggagg 240
ggaactcccc cggcccaagg ctgcgtgggt cggggtcgcg cggccgcaga aggggcgggg 300
tccgcccgcg aggggaggcg ccccccggga cccgagaggg gggtagggac cgcgggctgc 360
tggtgccggc gccggcagcgt gtgcggcggc caggggaggg cccgccccccg tcccgccccg 420
25 gctgcgagga ggaggcgccg cggccgcagg aggatgtact tggtgccggg ggacagggggg 480
ttggccggct cggggcacct ctcgtctcg ctgcgtgggt tgctgtctgt gccggcgcgc 540
tccggcaccc gggcgctggt ctgcctgcct tgtagcaggt ccaagtgcga ggagcccaagg 600
aaccggcccg ggagcatcgt gcaggcgcgt tgccgctgtc gtacacgtg cgccagccag 660
ggyaacgaga gctgcggcgg cacccgcggg atttacggaa cctgcgaccc gggctgcgt 720
30 tgtgtcatcc gccccccgct caatggcgcac tccctcaccg agtacgaagc gggcgttgc 780
gaagatgaga actggactga tgaccaaactg ctgggtttt aaccatgcaaa taaaacacctt 840
attgctggct gcaatataat caatgggaaa tgtgaatgtt acaccattcg aacctgcagc 900
aatcccttt agtttccaag tcaggatatg tgcccttcag ctttaaagag aattgaagaa 960
gagaagccag attgctccaa gccccctgt gaagtccagt tctctccacg ttgtcctgaa 1020
35 gattctgttc tgatcgaggg ttatgcicct cctggggagt gctgtccctt acccagccgc 1080
tgcgtgtgca acccccgcagg ctgtctgcgc aaagtctgcc agccggggaa cctgaacata 1140
ctagtgtcaa aagccctcagg gaagccggga gagtgcgtgtt acctctatga gtcaaaacca 1200
gttttcggcg tggactgcag gactgtggaa tgccctactg ttccgcagac cgcgtgtccc 1260
ccggcagct atgaaactca agtcagacta actcgagatg ttccgcgtac ttggccaacca 1320
40 agatgcgagt gtctctctgg cttagtgcgtt ttccgcgtgtt gtggatgtgg atccactccc 1380
cgcatagtc ctgcgtggca tgggacacact ggaaatgtct gtgatgtctt tgaatgtgtt 1440
aatgatacaa agccagcctg cgtatataac aatgtggaa attatgtatgg agacatgttt 1500
cgaatggaca actgtcggtt ctgtcgatgc caagggggcg ttgcctatcg cttcacccgc 1560
cagtgtggc agataaaactg cgagaggatc tacgtgcgc aaggagatg ctgcggcgt 1620
45 tgtgaagatc cagtgtatcc tttataataat cccgcgtggct gctatgcca tggcctgtatc 1680
cttgcacccacg gagaccgggt gcgggaaagac gactgcacat tctgcgtatc cgtcaacccgt 1740
gaacgcacact gcgttgcgac cgtctgcgca cagacctgca caaaccctgt gaaagtgcct 1800
ggggaggtttt gcccgtgtg cgaagaacca accatcatca cagttgatcc acctgcatgt 1860
50 ggggagttat caaactgcac tctgcacacgg aaggactgca ttaatggtt caaacgcgtat 1920
cacaatgggtt gtccggacactg tcagtgcata aacacccagg aactatgttc agaacgtaaa 1980
caaggctgca ccttgaaactg tcccttcgtt tccttactg atgccccaaa ctgtgagatc 2040
tgtgagtgcc gccccaaaggcc caagaagtgc agaccataa tctgtgacaa gtattgtcca 2100
cttggattgc tgaagaataat gcacggctgt gacatctgtc gctgtaaagaa atgtccagag 2160
55 ctctcatgca gtaagatctg cccctgggt ttccagcagg acagtacccgg ctgtcttatac 2220
tgcaagtgcg gagaggcctc tgcttcagct gggccaccca tcctgtcggt cacttgtctc 2280
accgtggatg gtcatcatca taaaaatgag gagagctggc acgtgggtt cggggaaatgc 2340
tactgtctca atggacggga aatgtgtgcc ctgcattcacct gcccggtgcc tgcctgtggc 2400
aaccggccacca ttcacccctgg acagtgcgtc ccatcatgtg cagatgactt tgggtgcag 2460
60 aagccagagc tcagttactcc ctccatgtc cacccccctg gaggagaata ctttggaa 2520
ggagaaacgt ggaacattga ctccgtact cagtgcaccc gcccacagcgg acgggtgtcg 2580
tgtgagacag aggtgtgccc accgcgtgcct tgccagaacc cctcagcgcac ccagattcc 2640
tgctgccccac agtgcatacaga tcaaccccttt cggccatttc tgcctgtggc tgcctgtggc 2700
ctaattact gcaaaaaatgat tgaagggat atattccgtt cagctgatgc ctggaaagcct 2760
gacgtttgcata ccagctgcat ctgcattgtat agcgttaatta gctgttttc tgcgttgc 2820
65 cttctgtat cctgtgaaag acctgtctgt agaaaaaggcc agtggttgtcc ctactgcata 2880
aaagacaccaa ttccaaagaa ggtgggtgc cacttcagtg ggaaggccctt tgccgacag 2940

65 mylvagdrgl agcghllvsl lgl111pars gtralvclpc deskceeprn rpgsivqgvc 60
gccytcasqq nescggtfgi ygtcdrglrc virpplngds lteyeagvce denwtddqll 120
gfkpcnenli agcniingkc ecntirtcsn pfefpsqdmc lsalkrieee kpdcskarce 180
vqfsprcped svlieqvapp qeccplpsrc vcnpaqclrk vcqpgnlhil vskasqkpqe 240

ccdlyeckpv fgvd crtvec ptvqqtacpp dsyetqvrlt adgcctlptr ceclsglcgf 300
 pvcevgstpr ivsrgdgtg kccdvfecvn dtkpacvfnn veyydgdmfr mdnrcrfcrcq 360
 ggvaicftaq cgeinceryy vpegeccpvc edpvypfnnp agcyanglil ahgdrwredd 420
 ctfcqcvnge rhcavatvcgq tctnpvkvpq eccpvceupt iitvdppacg elsncnltrk 480
 5 dcincfkrdh ngcrtcqcin tqelcserkq gctlncpfgf ltdaqnceic ecrprpkkr 540
 piicdkycpl gllknkhgcd icrckkcpel scskicplgf qqdshgclic kcreasasag 600
 ppilsgtclt vdghhhknee swhdgcrecy clngremcal itcpvpacgn ptihpgqccp 660
 scaddfvvqk pelstpsich apggeyfveg etwnidsctq ctchsgrvlc etevcppllc 720
 10 qnpsrtqdsc cpqctdqpf r pslsrnnsvp nyckndegdi flaaeswkpd vctscicids 780
 viscfsescp svscerpvlr kgqccpycik dtipkkvvch fsgkayadee rwldldscth 840
 yclqqqtcls tvscpplpvc epinvegssc pmcpemvpe ptnipiektn hrgevdlevp 900
 lwptpsendi vhlpdmghl qvdyrdnrlh psedssldsi asvvvpiiic lsiliaflfi 960
 nqkkqwipll cwyrtptkps slnnqlvsd ckkgrvqvq ssqrmiriae pdarfsgfys 1020
 mqkqnhlqad nfyqtv

15

<210> 35

<211> 716

<212> DNA

20 <213> Human

<400> 35

25 gcagtacctg gagtgtcctg cagggggaaa gcgAACCGGG ccctgaagt cggggcagtc 60
 accccggggct cctggccgc tctccggggc tgggcttag cagcgtatct gcttgcgtccc 120
 agaagtccag agggatcagc cccagaacac accctctcc ccgggacgccc gcacgtttct 180
 ggaggcttaga gaaaggcatga agagtgggtt ccacctgtgt gcccactgag aaaagaattt 240
 ccagaactcg gtcctatattt acagatttag aaactatggt tcaagaagag aggacggggc 300
 30 ttgagggaat ctcctgattc tcccttatatg acctcaaact gaccatacta aacagtgtag 360
 aaggctttt taaggctcta aatgtcaggg tctcccatcc cctgatgcct gacttgtaca 420
 gtcagtgtgg agtagacggg ttccctccacc cagggttgac tcagggggat gatctgggtc 480
 ccattctgtt cttaagaccc caaacaaggg tttttcagc tccaggatct ggacgcctcta 540
 35 tctggtagt gtcgtaacct ctgtgtgcct cccgttaccc catctgtcca gtgagcttag 600
 cccccatcca cctaacadagg tggcccacagg gattacttag ggttaagacc ttagaactgg 660
 gtctagcacc cgataagagc tcaataaaatg ttgttcctt ccacatcaaa aaaaaaa

<210> 36
 <211> 395
 40 <212> DNA
 <213> Human

<400> 36

45 ccaatacttc atttttcatt ggtggagaag attgttagact tctaaggcatt ttccaaataa 60
 aaaagctatg atttgatttc caacttttaa acattgcatt tcctttgcct tttactacat 120
 tctccaaaaa aaccttgaaa tgaagaaggc cacccttaaa atacttcaga ggctgaaaat 180
 atgattattt catttgcattt cttagccta tgttatattt cttaactttt gcactttcac 240
 50 gcccagtaaa accaaagtca gggtaaccaa tgtcattttt caaatgtta aaaccctaata 300
 tgcagttcct tttttaaattt attttaaaga ttacttaaca acatttagaca gtgcaaaaaaa 360
 agaagcaagg aaagcattct taattctacc atcct

<210> 37
 <211> 134
 55 <212> DNA
 <213> Human

<400> 37

60 ccctcgagcg gccggccggg caggtacttt taccaccgaa ttgttcactt gactttaaga 60
 aaccctataaa gctgcctggc tttcagcaac aggcctatca acaccatggt gagtctccat 120
 aagggacacc gtgt

<210> 38
 <211> 644
 65 <212> DNA

<210> 41
<211> 987
<212> DNA
<213> Human

5
<400> 41

aacagagact ggcacaggac ctcttcattg caggaagatg gtagtgttagg caggtAACAT 60
tgagctctt tcaaaaaagg agagctctc ttcaagataa ggaagtggta gttatgggtgg 120
10 taaccccccgg ctatcaGTCc ggatgggtgc caccCCTCCT gctgttagat ggaAGCAGCC 180
atggagtggg agggaggcgc aataagacac ccctccacAG agcttggcat catggaaAGC 240
tggTCTacc tcttcctggc tcctttgtt aaaggcctgg ctgggagcct tcctttggg 300
tgtCTTCTC ttcTCCAACC aacagaaaAG actgtCTTC aaaggTGGAG ggtCTTcatG 360
15 aaACACAGCT gCcAGGAGCC caggcacAG gCTGGGGCC tggAAAAGG AGGGCACACA 420
ggaggAGGGA ggAGGTGGTA gggAGATGCT gCTTTACCT aaggTCTCGA aacaAGGAGG 480
gcagaatAGG cagaggcCTC tccgtccccAG gCCATTttt gacAGATGGC gggACGGAAA 540
tgcaatAGAC cagCTGCAA gaaAGACATG tgTTTGTG acaggcAGTG tggCCGGGTG 600
gaacaAGCAC aggCCttggA atccAAAtGGA ctGAATCAGA accCTAGGCC tGCCATCTGT 660
20 cAGCCGGGTG acCTGGGTCA atTTTGTCT ctaAAAGCCT cAGTCTCCT atCTGAAAA 720
tgaggCTTGT gatacCTGTT ttGAAGGGTT gCTGAGAAA ttaAGATAA gggTATCCAA 780
aatAGTCTAC ggCCATACCA CCCTGAACGT GCCTAACTCt GtaAGCTAAAG CAGGGTCAGG 840
CCTGGTTAGT acCTGGATGG ggAGAGTATG gaaaACATAc CTGCCCGCAG ttggAGTTGG 900
actCTGTCTT AACAGTAGCG tggcacACAG aaggCACTCA gtaAAataCTT gttGAATAAA 960
tgaagtAGCG atttggTGTG aaaaaaaaa

25
<210> 42
<211> 956
<212> DNA
<213> Human

30
<400> 42

cggacggTgg ggcggacGCG tgggtgcagg agcaggGCgg ctGCCGACTG ccccaACCAA 60
ggaaggAGCC cctgAGTCCG CCTGCGCCTC catCCATCTG tccggCCAGA gCCGGCATCC 120
35 ttgcCTGTCT aaAGCCTAA ctaAGACTCC CGCCCCGGGC tggCCCTGTG cAGACCTTAC 180
tcAGGGGATG ttTAACCTGGT gCTCGGGAAg ggAGGGGAAG gggCCGGGGa gggGGCACGG 240
caggCgtGTG gcAGGCCACAC gCAGGCGGCC AGGGCGGCC gggACCCAAA gCAGGATGAC 300
caCgCACCTC caCgCCACTG CTCCTCCCA atGATTtGG aACCAAAAGTC tAAActGAGC 360
40 tcgCAGCCCC CGCAGCCCTCC CTCGCGCTCC catCCCGCTT agCgCTCTGG acAGATGGAC 420
gcaggCCCTG tCCAGCCCC AGTGCCTCG ttCCGGTCCC caCAGACTCG CCCAGCCAAc 480
gagATTGCTG gaaACCAAGT caggCCAGt gggCGACAA aaggGCCAGG tgcGGCCtGG 540
ggggAACGGA tgCTCCGAGG ACTGGACTGT ttTTTCAcA catCGTTGCC gCAGCGGTGG 600
gaaggAAAGG cAGATGAAA tgATGTGTTG ttTACAGGG tatATTTTG atACCTCAA 660
tgaattaATT cAGATGTTT ACgCAAGGAA ggACTTACCC AGTATTACTG ctGCTGTGCT 720
45 ttGATCTCT GCTTACCGTT CAAGAGGCGT gtGcAGGCCG AGTCTGGT ACCCCATCAC 780
tcgCAGGACC AAGGGGGCGG ggACTGCTGG CTCACGCCc GCTGTGTCT CCCTCCCTC 840
CCTTCCTTGG gcAGAAATGAA ttCGATGCGT ATTCTGTGGC CGCCATCTC gCAGGGTGGT 900
gttattCTGT cATTACACa CGTCGTTCTA attAAAAAGC gaAttATACT ccaAAA

50 <210> 43
<211> 536
<212> DNA
<213> Human

55 <400> 43

aaATAAAACAC ttCCATAACA ttttGTTTtC gaAGTCTATT aATGCAATCC cACTTTTtC 60
cccCTAGTTt ctaAAATGTTA aAGAGAGGGG AAAAAGGCT cAGGATAGtT tTCACCTCAC 120
agtGTTAGt GTCTTTTATT ttACTCTTGG AAATAGAGAC tCCATTAGGG ttttGACATT 180
60 ttGggGAACCC agTTTACCA ttGtGTCAGt AAAACAATAA gATAGTTGA gagCATATGA 240
tctAAATAAA gACATTGAA gggTTAGTT GAATTCTAA AGTAGGTAAT AGCCAAATAG 300
cattCTCATC CCTTAACAGA CAAAACCTA tttGtCAAA GAATTAGAAA AGGTGAAAAT 360
atTTTTTCCA gATGAAACTT gtGCCACTC CAATTGACTA ATGAAATAcA AGGAGACAGA 420
ctggAAAAG TGGGTTATGC cacCTTAAAC ACCCTTCTG gtaAAATATTA TGGTAGCTAA 480
65 agggTGGTTT cCCCGCACC TGGACCTGGA CAGGTAGGGT tCCGTGGTTA ACCAGT

<210> 44
 <211> 1630
 <212> DNA
 <213> Human

<400> 44

```

ggggaggggac gagtatggaa ccctgaaggt agcaagtcca ggcactggcc tgaccatccg 60
gctccctggg caccaagtcc cagggcaggag cagctgtttt ccatcccttc ccagacaagc 120
tctattttta tcacaatgac cttagagag gtctcccagg ccagctcaag gtgtcccact 180
atccccccttg gagggaagag gcaggaaaaat tctccccggg tccctgtcat gctactttt 240
ccatcccttgt tcagactgtc caggacatct tatctgcagc cataagagaa ttataagga 300
gtgattttcc tttaggcccag gacttgggcc tccagctcat ctgttccctc tgggcccatt 360
catggcagggt tctgggctca aagctgaact ggggagagaaa gagatacaga gctaccatgt 420
gactttacct gattgccctc agtttggggt tgcttattgg gaaagagaga gacaaagagt 480
tacttggttac gggaaatatg aaaagcatgg ccaggatgca tagaggagat tctagcaggg 540
gacaggattg gctcagatga cccctgaggg ctcttccagt ctgaaatgc attccatgtat 600
attaggaagt cgggggtggg tgggtgttgg gggctagttt gggttgaatt taggggcccga 660
tgagcttggg tacgtgagca ggggtgttaag ttaggtictg cctgtatttc tgggtcccctt 720
ggaaatgtcc ccttcttcag tgtcagacct cagtccctgt gtccatatcg tgcccagaaa 780
agtagacatt atccgtcccc atcccttccc cagtgcactc tgaccatgtc agtgcctgt 840
gcccagtgtac ctgggggagc ctggctgcag gcctctactg gtcccttaaa cttgggtg 900
tgtgattcag gtcccccaggg gggacttcagg gaggaaatatg gctgagttct gtatgttcca 960
gagttggctg gtagagccctt ctagaggttc aaaaatattttt accaaagccc ctacctgt 1020
atggaaattgg ctgaggatca aacgtatgtt ggtgaaagga taccaggatg ttgtctaaagg 1080
tgagggacacat tttgggtttt ggacttacca gggtgtatgtt agatctggaa cccccaagtg 1140
aggctgggagg ggtttaaggt cagttatggaa gatagggttg ggacagggtg ctttggaaatg 1200
aaagagtgtac cttagagggc tccttgggcc tcaggaatgc tcctgctgt gtgaagatga 1260
gaaggtgctc ttactcagtt aatgtatgtt gactatattt accaaagccc ctacctgt 1320
ctgggtccct ttagcacag gagactgggg ctaaggccc ctcccaggga agggacacca 1380
tcaggcctct ggctgaggca gtagcataga ggatccattt ctacctgcat ttcccagagg 1440
actagcagga ggcagccctt agaaaaccggc agttcccaag ccagccctg gctgttct 1500
cattgtcact gcctctccc caacctctcc tctaaccac tagagattgc ctgtgtccctg 1560
cctctgtccctt ctgttagaat gcagctctgg ccctcaataaa atgcttctgtt cattcatctg 1620
aaaaaaaaaaa
  
```

<210> 45
 <211> 169
 <212> DNA
 <213> Human

<400> 45

```

tcttttgcctt ttagctttttt atttttgtat taacaggagt cttattacac ataggctctga 60
taaaaacttgtt ttatgtatctt cagtcgttcc ccaagtcgtc ataacttagat aacgtatgaa 120
ggaaaaaacga cgacgaacaa aaaagtaagt gtttggaaaga ctttagttga
  
```

<210> 46
 <211> 769
 <212> DNA
 <213> Human

<400> 46

```

tgcagggtcat atttactatc ggcaataaaaa ggaagcaaaag cagtattaag cagcgggtgga 60
atttgtcgct ttactttttt ataaaagtgtc acataaaaatg tcataatttcc aaattttaaaa 120
acataactcc agttcttacc atgagaacag catggtgatc acgaaggatc ttcttggaaa 180
aaacaaaaaac aaaaacaaaaa aacaatgtc tcttctgggt atcacatcaa atgagataca 240
aagggtgtact aggcaatctt agagatctgg caacttattt tataatataag gcatctgt 300
ccaagagacg ttatgaatta aatgtacaaa tttttttttt ataaatgtat taaaatgtcaag 360
cttcatataaa tgacaccaat gtctcttaagt tgctcagaga ttttggggctt ctgtggccct 420
ggccagtcctt tttctgtata gtctgttcc gccttcataat ataggcagct cctgtatcc 480
catggccatgt aatggggaaa caagcatgga atataaaaac tttaacatggaa aactttgtt 540
tattttgtaa taaaatctt tttcccttgg aaaccttcaaa aactttgttca gaatgttggg 600
ttgatataatg tgacaatgtt gtaccttctt agtgcacatggaa aacatcattttt tttctgtctg 660
cctgcctttttt tttttttttt aatgtttttt atcattttttt caaattttttt tttttttttt 720
  
```

ggacatgtt acggagagga aaggttaggaa agggtttaggg atagaagcc

<210> 47

2529

<212> DNA

1400> 47

| | |
|----|--|
| 10 | tttagttcat agtaatgtaa aaccatttgt ttaattctaa atcaaatcac tttcacaaca 60 gtgaaaatta gtgactgggt aagggtgtcc actgtacata tcatcatttt ctgactgggg 120 tcaggacctg gtcctagtcc acaagggtgg caggaggagg gtggaggcta agaacacaga 180 aaacacacaaa aagaaggaa agctgcctt gcagaaggat gaggtggta gcttgcgag 240 ggatgggtgg aagggggctc cctgttgggg ccgagccagg agtcccaagt cagctctcct 300 gccttactta gctccctggca gagggtgagt ggggacctac gaggttcaaa atcaaatggc 360 atttggccag cctggcttta ctaacaggtt cccagagtgc ctctgttgc tgagctctcc 420 tgggctcaact ccatttcatt gaagagtcca aatgattcat ttccttaccc acaactttc 480 attattcttc tggaaaccca tttctgttga gtccatctga ctaagtctct ctctcttagag aagagacact 600 actagttggg gccactgcac tgaggggggt cccaccaatt ctctcttagag aagagacact 660 ccagaggccc ctgcaactt gcggatttcc agaagggtat aaaaagagca ctcttgagt 720 ggtgc当地 cggcatatga accacactt caaacttta tattttgtct gtttaaacac 780 tgaactctgg tggacagg tacaaaggag aagagatggg gactgtgaag aggggaggggc 840 ttccctcattt ttcctcaaga tctttgttcc catabactt gcaacttgc ttgagaaaaaa 900 gcaatagatg gggcttccat ccattttgtt gttattgtct gggtagcca ggacactgt 960 ggatggcaaa gtaggagaga gcccagagg aaaggccatc tccctccagc tttgggtt 1020 ccagaaagag gctggatttc tggatgaag cctagaaggc agagcaagaa ctgttccacc 1080 aggtgaacag tcctacctgc ttggtaccat agtccctcaa taagattcag aggaagaagc 1140 ttatgaaact gaaaatcaaa tcaaggattt gggagaata atttccccctc gattccacag 1200 gagggaaagac cacacaatat cattgtgtc gggctccccca aggccctgcc acctggctt 1260 acaaatcatc aggggttgcc tgcttggcag tcacatgtt ccctggttt agcacacata 1320 caaggagttt tcagggact ctatcaagcc atacaaaat cagggtcaca tggggtttc 1380 cccttccttgc ctgtttcat aaaagacaac ttggcttctg aggtatgggtt tctttgcatt 1440 gcagttgggc tgacctgaca aagccccccag ttccctgttgg caggttctgg gagaggatgc 1500 attcaagtt ctgcagccata ggggacagggt ctgttgcattt agtttattact gcctcgagc 1560 tccaaatccc accaaagtcc tgactccagg tctttcttgc tgcacagttag tcagtcctcag 1620 cttcggcagt attctcggtt gtatgttctc tggcagagag aggcatgtt acatagttt 1680 agggagaaag ctgatggaa acctgtgagt taaggccatc gtctcaccag gaataattta 1740 tgccaggaaa ccaggaagtc attcaagttt ttctctgagg caaaagacac tgagcacagc 1800 ccagagccaa taaaagatct ttgagtctct ggtgaattca cgaagtgcacc ccagctttag 1860 ctactgcaat tatgatttt atgggacagc aatttcttgc atctctacag aggaagaaga 1920 gggggagttt gaggggaaagg aaagagaaca gagcggcact gggatttggaa aggggaacct 1980 ctctatctga ggagccccca ctggcttctg aagcaactt ocaaggggtt tttaaagaca 2040 tgaaaatttc cagaaatacc atttggtgc tcccttgcatt tctgtaatat taaactcagg 2100 tgaaaattata ctctgacagt ttctctctt ctgcctcttgc cctctgcaga gtcaggacat 2160 gcagaactgg ctgaaacaaag atttcatgtt gtcacccatg agagatgact caatgccaag 2220 gcctgaaatgtt atagatgtt tacagcggtt gcgatattca ggggttcatcg ccaactggtc 2280 tcgatgttca aagctctgtt gaagaaacaa gactcttgc ttttttttttttctg atcccaactgta 2340 ttccaggagt caagatttgc caggaagcca aacaccaggat gtttttttttttctg cactgtccacca 2400 gtccagagcc ctgcccacggc tgtacgcagg agcccgatcat taggcaatc ggagccagaa 2460 catgtatcacc agggccacaa ataggaagag gcgtgacagg aactgctcgtt ccacatactt 2520 qqqqtgtcc |
| 15 | |
| 20 | |
| 25 | |
| 30 | |
| 35 | |
| 40 | |
| 45 | |
| 50 | |

<210> 48

<211> 1553

<212> DNA

<213> Human

<400> 48

```

60 tttttttttt tttttgattt ctgggacaaat taagcttat ttttcatata tatatatattt 60
      ttcatataat tataacata catataaaa ggaaacaaatt tgcaaatttt cacacctgac 120
      aaaacccatata atacacacat atgtatgcattt acacacagac agacacacac acccgaagtt 180
      ctagccaggc ccgttttcca tccctaaatgat ccattctctc atttgggccc ttctagggtt 240
      ggggcctgat gcttggtttt tagaagttt gtgctaataat aaccatagct ttaatccccca 300
      tgaaggacacat tttttttttt atctttgtct gctccccgct gcctttcaatgat tttacgtgat 360

```

ccatcaagag ggctatgggaa gccaagtgaac cacggggat tgaggctaatt tcacctgaac 420
 tcgaaaacag cgcccgactt cctcaccgcgaa ggcacgcgtc ttttctttt ttttcctcgaa 480
 gacggagttct cgctgtgttgc cccaggctgg agtgcagtgg cacggtctcg gctcaactgca 540
 5 agctccacact cctggattca taccattctc ctgcttcagc cttccggatgta gctgggacta 600
 taggtgccaa ccactacgccc tagctaattttttttagt tttttgtat ttttagtaga gacagggttt 660
 caccgtgttgc gccaggatgg tctcgccctg actttgtat ccggccggct cggcctccca 720
 aagtgcgtggg attacaggcg tgagccacca cacctggccc cggcacgtat ctttaagga 780
 atgacaccag ttccctggctt ctgaccaaag aaaaaatgtc acaggagact ttgaagaggc 840
 10 agacaggagg gtggtggcag caaacactgca gctgcttcgt gatgctgctg ggggtgtcctc 900
 cggagcgggt gtgaacacagcg cactcaaca tgagcaggcg cctggctccg gtgtgtcctc 960
 acttcagttgc tgccacccgttga tgggtggaaac cagcctttgg ggcaggaaac cagtcagag 1020
 aggctaccca gctcagtc tggcaggaggc caggatattta cagccataat gtgtgtaaag 1080
 aaaaacacgc ttctgcaga aactctccca cccgctcggtt agactggggc tccttgcttgc 1140
 15 ggatgagttct cactaacatgttggatgggtt gttggactgtt ccctgaaaag cgggccttgc 1200
 agggccaaatgttggatggcataac ccagtggccc tctgaaaggg ggtgtgcagg 1260
 cgaggggggaggc aggagggttcc tctctagttcc ctttggaggc tttggctgag agaagagtga 1320
 gcagggggacttggatggcataac ccaggcaggga agggagctga agtgattcgg ggctaatgcc 1380
 tcagatcgat gtatttctctt ccctggcttc cccggagccctt cttgtcaccgc ctgtgcctc 1440
 20 gcaggaggccc catctcttctt gggagcttat ctgacttaac ttcaactaca agttcgctct 1500
 tacgagacccg ggggttagcgt gatctccgttccctgagc gcctgcacccg cag

 <210> 49
 <211> 921
 <212> DNA
 25 <213> Human

 <400> 49

 ctgtggtccc agctactcag gaggctgagg cgggaggatt gcttggagccc aggagttgg 60
 30 tgttgcagtgc accaagatc gcaccatttc cctccactctt gggccacggc gcaataaccct 120
 gtctcagaaa acaaacaaca aaaagcagaa acgtgttgcgg ggtcggttta cgggaaaacc 180
 gcctgtcaga acacttggctt actccttccca cagatcagtgc gacctggaa tgagggttgg 240
 tccccggagg cttttcttccca agtctgttgc accagaccggc ccattggaaac cctggccaca 300
 gaaggcctccc ggggagtgttgc ccagacccgtt gaccgtgttgc ctgtgtgtc tgggtgtgg 360
 35 ggagggtggg ggtgtgttgc aaagggtgttgc gttggccgggg ggtgttcatg ggcaagcatg 420
 tgcgtgcctt tgggtgttgc tggcccttccca ctgcagccgtt cgggtgttgc tccctccac 480
 cccttcgcca ccttctgagc attgtctgtc cacgtgagac tggccagaga cagcagagct 540
 ccacgtgtttttaagggggaccccttccca ggacccgttgggg gtcctgcgtt atctccatgac 600
 cagggtctaa atgacccgac atgcattacc tggcccttccca tgaccacactt ccctgtcccc 660
 40 gtccctgcgttccca cctggccccc tggcgtctca cgggtgttgc tggccctgttgc attgggtttc 720
 actgttagcaa actacattctt ggtggaaat tttcatgttac atgtgtggca tggaaaat 780
 ttcaaaataaa atggacttga tttagaaaagc caaaaagctg tgggtccctt ccagcacgg 840
 tactttgacc tcttgccttccca aacccttccca ttgggtccga ggctggtagc tttgttcaact 900
 tcagatgggtt gggggcgggtt g

 45 <210> 50
 <211> 338
 <212> DNA
 <213> Human
 50 <400> 50

 atgatcttac tagatgcctt accgtaaaaat caaaaacacaa aacccttactg actcattccc 60
 tcccttccag atatttccccc atttctcttac ttcccttccatgtt agccaaactt tccaaaaattt 120
 55 catgttttgtt ctttcttccca tcatgttcaaa cccaccctgtt ctttagctacc acccccttgcgt 180
 aacgacccatgttccca cctgggttgc aacaaatgtc agcatgatac cataactcaat gatcccttgcgt 240
 cactgttgcgttccca attgtcatca ttccatggcc ttactttccca tctcagcggcc atttgcata 300
 gtaagaaaacttccca ttctttcttgcgtt aatttttttttgcgtt tcttttttgcgtt

 60 <210> 51
 <211> 1191
 <212> DNA
 <213> Human
 65 <400> 51

| | | | | | | |
|--------------|-------------|-------------|-------------|-------------|-------------|------|
| cttagcaagca | ggtaaacgag | ctttgtacaa | acacacacaa | accaacacat | ccggggatgg | 60 |
| ctgtgttgttgc | ctagagcaga | ggctgattaa | acactcagtgc | tgttggctct | ctgtgccact | 120 |
| cctggaaaat | aatgaattgg | gtaaggaaaca | gtaataaga | aatgtgcct | tgctaactgt | 180 |
| gcacattaca | acaaagagct | ggcagctcct | gaaggaaaag | ggcttgcgc | gctgccgttc | 240 |
| aaacttgtca | gtcaactcat | gccagcagcc | tcagcgctcg | cctccccagc | acaccctcat | 300 |
| tacatgtgtc | tgtctggcct | gatctgtgc | tctgctcgga | gacgctcctg | acaagtgcggg | 360 |
| aatttctcta | tttctccact | ggtgcaaaga | goggatttct | ccctgcttct | cttctgtcac | 420 |
| ccccgtcct | ctccccccagg | aggctccttg | atttatggta | gctttggact | tgcttccccg | 480 |
| tctgactgtc | cttgacttct | agaatggaag | aagctgagct | ggtaaggga | agactccagg | 540 |
| ccatcacaga | taaaagaaaa | atacaggaag | aaatctcaca | gaagcgtctg | aaaatagagg | 600 |
| aagacaaaact | aaagcaccag | catttgaaga | aaaaggcctt | gagggagaaa | tggctcttag | 660 |
| atggaatcag | cagcggaaaa | gaacaggaag | agatgaaagaa | gaaaatcaa | caagaccagc | 720 |
| accagatcca | ggttctgaa | caaagtatcc | toaggcttgc | gaaagagatc | caagatcttg | 780 |
| aaaaagctga | actgcaaatc | tcaacgaaagg | aagaggccat | tttaaagaaaa | ctaaagtcaa | 840 |
| ttgagcggac | aacagaagac | attataagat | ctgtgaaagt | ggaaagagaa | gaaagagcag | 900 |
| aagagtcaat | tgaggacatc | tatgctaata | tccctgacct | tccaaagtcc | tacataccct | 960 |
| ctaggttaag | gaaggagata | aatgaagaaa | aagaagatga | tgaacaaaat | aggaaagctt | 1020 |
| tatatgccat | gaaaattaaa | gttggaaaaag | acttgaagac | tggagaaaagt | acagttctgt | 1080 |
| cttccaatac | ctctggccat | cagatgactt | taaaaggtac | aggagaaaa | gtttaagatg | 1140 |
| atgggcaaaa | gtccagtgt | ttcagtaaag | tgctaatcac | aagttggagg | t | |

<210> 52
<211> 1200
<212> DNA
25 <213> Human

<400> 52

| | |
|----|--|
| 30 | aacaggggact ctcactctat caaccccagg ctggagtcgg gtgcgc(cc)ac cctggctccc 60 tgcaacccctcc gcctccccagg ctcaga(ac)ac tctcc(tg)ctt cagtgc(t)ct agtagctggg 120 actacaggca cacaccacca tgcccagcca atttttgcat tttttgtaga gacagggttt 180 cgccttctgt ccaggccggc atcatataact ttaaaatcatg cccagatgac ttaataacct 240 aataacaatat atcaggttgg tttaaaata attgctttt tattat(tttt) gcattttgc 300 35 accaaccccta atgctatgta aatagtgtt atactgttgc ttaacaacag tatgacaatt 360 ttggctttttt ctttgttata ttttgttattt tttttttttt ttgtgtggc tttttttttt 420 ttctcagtgt tttcaattcc tccttgggtt aatccatgga tgcaaaaccc acagatatga 480 agggctggct atatatgcat tgatgattgt cctattatata tagttataaa gtgtcattta 540 atatgttagt aaagttagtgg tacagtggaa agagttagttt aaaaacataaa catttggacc 600 tttcaagaaa ggtagcttgg tgaagttttt cacc(t)caa(c) ctatgtccca gtcagggctc 660 40 tgctactaat tagctataat ctttgaccaa attacatcac ctttggactt cagttgcctc 720 acctgtaaaaa tggaaagaact ggatacttc taagg(t)cact tccagccccgt tcatttata 780 actctgttat gctgaggaag aaattcacat tgg(t)taact gtatgagtca aactgaaaat 840 gattattaaa gtggggaaaaa gccaattgtt cttcttagaa agctcaacta aatttggagaa 900 45 gaataatctt ttcaattttt taagaattt aatattttt aagggttgc ac tttttttttt 960 agagatgggg tctcactctg tcacccagac tggagtacag tggcacaatc atagctact 1020 gctgcctcaa attcatgggc tcaagtgtac tctcc(tg)ctc tggcctccaga gtagctgcga 1080 ctatgggcat gtgccaccac gcctggctaa cattttgtattt gacctattta tttattgtga 1140 tttatatctt tttttttttt tttttttttt ttttttacaa aatcagaaaat acttattttg 1200 |
|----|--|

50 <210> 53
<211> 989
<212> DNA
<213> Human

55 <400> 53

| | | | | | | | |
|----|-------------|-------------|------------|-------------|-------------|------------|-----|
| | aagccaccac | tcaaaaacttc | ctatacattt | tcacagcaga | gacaagtgaa | catttatttt | 60 |
| | tatgccttc | ttccttatgtg | tatttcaagt | cttttcaaa | acaaggcccc | aggactctcc | 120 |
| | gattcaatta | gtccttggc | tggtcgactg | tgcaggagtc | cagggagcct | ctacaaatgc | 180 |
| 60 | agagtgactc | tttaccaaca | taaaccctag | atacatgcaa | aaagcaggac | ccttcctcca | 240 |
| | ggaatgtgcc | atttcagatg | cacagcaccc | atgcagaaaa | gctggaaattt | tccttggAAC | 300 |
| | cgactgtgat | agaggtgctt | acatgaacat | tgctactgtc | tttctttttt | tttgagacag | 360 |
| | gtttcgcttg | tgcccaggct | gagtgcaatg | cgtgatctca | ctcaactgcaa | ttccacacct | 420 |
| 65 | aggttcaagc | attctcctgc | tcagcctcct | agttagctggg | ttacaggcac | tgccaccatg | 480 |
| | ccggctaatt | ttgtatTTT | gtagagatgg | attttctccat | ttggtcaggc | ggtctcgaac | 540 |
| | cccaacaccta | gtgatctgcc | acctcagcct | cctaagtgtt | ggattacagg | atgagccacc | 600 |

aaaatgtaaa cttcacctag ttcatcttct ccaaataccca agatgtgacc ggaaaagtag 420
 cctctacagg acccactagt gcccacacag agtggttttt ctgcacactg ctgttcaca 480
 ggactttgtt ggagagtttag gaaattccca ttacgatctc caaacacgta gttccatac 540
 5 aatcttctg actggcagcc ccggatataca aatccaccaa ccaaaggacc attactaat 600
 ggcttgaatt ctaaaagtga tggctcact tcataatctt tcccccttat tatctgtaga 660
 attctggctg atgatctgtt tttccattg gagtctgaac acagtatcg taaattgtat 720
 tttatatcg tggatgtct atccacacgca catctgcctg gatctggag cccatgagca 780
 aacacttcgg ggggctgggt ggtgctgttg aagtgtgggt tgctccctgg tatgaataa 840
 10 ggcacgttgc acatgtctgt gtccacatcc agccgttagca ctgagcctgt gaaatcactt 900
 aaccatcca tttctccat atcatccagt gtaatcatcc catcaccaag aatgatgtac 960
 aaaaacccgt cagggccaaa gaggcgttgc cctccagat gcttctgtg gagttctgca 1020
 acttcaagaa agactctggc tttctcaa

<210> 59

15 <211> 747

<212> DNA

<213> Human

<400> 59

20 ttttcaaat cacatatggc ttctttgacc ccatcaaata actttattca cacaacgtc 60
 ccttaattt caaaggctca gtcattcata cacattaggg gatccacagt gttcaaggaa 120
 cttaaatata atgtatcata ccaacccaag taaaccaagt aaaaaaaaata ttcatataaa 180
 gttgttcaca cgtaggtcct agattaccag cttctgtca aaaaaaggaa atgaagaaaa 240
 25 atagatttt taacttagtat tggaaactaa ctttgtcct ggcttaaaaac ctccctcact 300
 ctcgtctgtc ccacacaaat gtttaagaag tcactgcaat gtactccccg gctctgtat 360
 aaagaagccc ctggcacaaa agattccagt gcccctgaag aggctccctt cctctgtgg 420
 gctctccctag aaaaccagcg ggacggccctc cctgtgtata ccgtctataa ctttaggggg 480
 30 ccctcgggca ggcaacggca gtggactcat ctcgggtatg gctgttagatg ctaacactgg 540
 ccaattcaat gccacaccta ctggttaccc tttgaggcga tttctccaga cagaagcccc 600
 ttgaaggccta ggttagggcag gatcagagat acacccgtgt ttgtctcgaa gggctccaca 660
 gcccagtacg acatgttgc agaagtagta tctctggact tctgcctcca gtcgaccggc 720
 cgccgaaatttta gtagtaatag cggccgc